SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	1	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence	:	
16223	46591	A	16318	219	1912	PHPTRTRSAQSVQYIEKDWGPG
10223	10371	` `	10510	217	1,512	EGLEEGGVLGQLVAFDSFTVSF
		ľ				HFVSAGNSPSDESEERERDPKV
						LTFPEYITSLSDSGTKHMAAGV
						RMECHSKGRCPSSCPLCHVTSS
		İ				PDTPAEPVLLEVTKAALLNTQS
		l				1
						FPCSLMQLLQEATMSSLWCSGT
						GDVIEDWCRCDSTAFGGCVPVS
		l				CSFLSAHRLRLSTVHEPSSTLVV
						LEWEHSEPPIGVQIVDYLLRQE
						KVTDRMDHSKVETGEHLLLHG
						LVLIQGDKTKTKKNALPHGFAP
		l				GPQYPRAMQEEQGEEEAVLLL
						QEAAVLLPLRVRDIRSCCFRHR
						ELLFLMLFPFLLLWAETVLSFV
						DDIISGAKSPCAMPSQVPDKQL
						TTIYFQPLIARFSFSSRKGLLRIM
						SSHCHGDCDVSTLDSYTVSAN
						HALPLPARFTLWGVDNTGRRS
						RPSDVIVYM*ITYFPFPPSSEIAD
						KIYNLFNGYTSGKEQQTAYNTL
						LDLGSPTLHRVLYHYNQHYESF
						ASF*LKYPLCLIHRKAGLILSQL
						GDLSSWCNGLLQEPKISLRRSS
		l				LKYLGCRYSEIKPYGLDWAELS
16224	46592	Α	16319	I	2031	
16225	46593	Α	16320	1	739	MKRAGSSGGRGECDISGAGRL
						GLEEAARLSCAVHTSPGGGRRP
						GQAAGMSAKERPKGKVIKDSV
						TLLPCFYFVELPILASSVVSLYF
						LELTDVFKPVHSGFSCYDRSLS
						MPYIEPTQEAIPFLMLLSLAFAG
						PAITIMVGEGILYCCLSKRRNG
						VGLEPNINAGGCNFNSFLRRAV
						RFVGVHVFGLCSTALITDIIQLY
						TGHQAPYFLTVRTPNYTSLNVS
						CT*NSYLAEDIRSRSDLTLTHSG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
16226	46594	A	16321	48	1359	GDKNQNPRNWPLGEGKGNSLG
		j				RKETGRSNSRSHSSRSKSR*QSS
						SRSRSRSHSRKKRHSSRSRSRTY
						SRSWSRDRMYSRDDRH*YEVIV
		l				DCSRHGREYKGREPRHGSGSLE
						RPTEQETSQPKIRKEIVETKGTA
						RGNCGTTGLLRPKTMMVERSY
						GPQDRVAVGTWPHPGTERWAS
						SWDVSSSVRHQSEGVVQGWEH
						RCTHKSQQEVGACVNFRTPKVI
						LIYNLYLPVSHQFSKTFMSIVIV
						QKNLYLHNDFRALNVTHIYRLT
			:			IPNTTTCFTATQDEDKQIFHTAA
						QHRSHTSSYQYLSLTRSHRYEI
						VFYTRGLGAPFPAGDGHGKFQ
						RLGKPYSHSVHFVIPQDKTQHIS
						INTPHKNSETTKQHTCSFAYGL
						YYYQQKEDNPYAWPQVPAAA
						EMVSRLAQSRHSQQSILIHFSHQ
						SYHTDHISTV
16227	46595	Α	16322	227	3020	SCIKPRRKEMGRSNSRSHSSRSK
						SRSQS\SSRSRSRSHSRKKRYSS\
		ŀ				RSR\SRTYSRSRSRDRMYSRDY
				ļ		RRDYRNNRGMRRPYGYRGRG
						RGV/YYQGGGGRYHRGGYRPV
						WNRRHSRSPRR\GRSRSRSPKR
						RSVSSQRSRSRSRRSYRSSRSPR
						SSSSRSSSPYSKSPVSKRRGSQE
						KQTKKAEGEPQEESPLKSKSQE
						EPKDTFEHDPSESIDEFNKSSAT
						SGDIWPGLSAYDNSPRSPHSPSP
16228	46596	В	16323	258	3728	IATPPSQSS
16229	46597	В	16324	1	435	
16230	46598	A	16325	1120	1209	ASSDNPP*SEPRVSLTIEGQEIDF
10230	40370	^	10323	1120	120)	FLDTGTAFSVLSSCPGWLSSRS
						VTIRGILGQPVTRYFSHLL
16231	46599	Α	16326	1291	1582	IHQSALCI*LKVCKHTNQQPVSS
10231	70377	, ,	10320	1271	1502	SGFVNAPIDTLYLATLVGPWRT
						LCVHTLYLVNLVGMWRTFVSS
						SGIVNAPISALSKQTTWLYQSA
						GCGWGQIRE
				L <u>.</u>	<u></u>	OCO MOČIVE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence	1	
16232	46600	Α	16327	12	840	APAVATEVLPGAGLTAGVLLW
			ŀ			SDPKEPSNPGSSHRVGASDLDN
					:	PAHPTLLSYTMQKLEEELDVVL
]					FDRSGHRTKFTNVGRMLLERG
						RVLLQAADKLTTDAEALARGW
						ETHLTIVTEALVPTPAFFPLIDK
						LAAKANTQLAIIT*VLAGAWER
						LEQGRADIVIAPDMHFRSSSEIN
						SRKLYTLMNVYVAAPDHPIHQ
						EPEPLSEVTRVKYRGIAVADTA
						RERPVLTVQLLDKQPRLTVSTIE
						DKRQALLAGLGVATMPYPMVE
	·					KDIAEGRLRVVSPESTS
16233	46601	A	16328	3	923	RFLSDGTIISVDSAGKVQFWDS
10233	40001	Γ	10326	3	1923	ATGTLVKSHLIANADVQSIAVA
						DFFPGHESRATEALCWAEGQR
						LFSAGLNGEIMEYDLQALNIKY
	,					AMDAFGGPIWSMAASPSGSQL
						LVGCEDGSVKLFQITPDKIQFER
						NFDRQKSRILSLSWHPSGTHIA
						1 -
1						AGSIDYISVFDVKSGSAVHKMI
						VDRQYMGVSKRKCIVWGVAFL SDGTIISVDSAGKVQFWDSATG
						1
						TLVKSHLIANADVQSIAVADQR RQFRGGHSRGNSLPFSAGPC\AS
						NSS*EAVGADKTVPASHS*RAH
į		ĺ	}			1
16234	46602	A	16329	2	1059	CGPTAQQR*YLGGTDTHLVFRP
16235	46603	Ā	16330	582	1540	
16236	46604	A	16331	1	720	
16237	46605	A	16332	1	477	
16238	46606	Ā	16333	1	540	
16239	46607	A	16334	1	261	
16240	46608	Ā	16335	964	1574	LRWCCGSQLFRGYWGGVTGR
						YLTVLPPHLAVGCMAPWGPFS
		ĺ		}		VRIAAVVFIQIFAGEVFWCPVM
						CVRGLCTPVLTGVPGHGNSHIP
		ł				AIVFTCSLQTLVGGTRRWWRC
						CAAGDVTKFPALVGVGGETPR
1						PQLPIGVTTVILLQVLVWKFGR
				1		NVSMSVWALRVPGLAFQSRH/
						ASWKFCRHSFLHCNAYIPLHTQ
				1		ECRSSHISFLDQ
16241	46609	Α	16336	3	87	DOMOSTITOT DDQ
16242	46610	A	16337	1	1398	
10272	1,0010	<u>'``</u>	1.0557	Γ,	1.570	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
16243	46611	Α	16338	1596	2205	LRWCCGSQLFRGYWGGVTGR
						YLTVLPPHLAVGCMAPWGPFS
						VRIAAVVFIQIFAGEVFWCPVM
						CVRGLCTPVLTGVPGNGNSHIP
						AIVFTCSLQTLVGGTRRWWRC
						CAAGDVAKFPALVGVGGETPR
						PQLPIGVTTVILLQVLVWKFGR
						NVSMSVWALRVPGLAFQSRH/
]			ASWKFCRHSFLHCNAYIPLHTO
						ECRSSHISFLDQ
16244	46612	В	16339	276	442	
16245	46613	A	16340	1	300	
16246	46614	В	16341	1	903	
16247	46615	Α	16342	119	889	RQHILESHSRPILCTTWLPLLIIQ
						DPRTLQSAAMLLTQSLFGGIFT
						RTRMKFGAVTWIGGPPLGDQS
						P\SSCSLLREKDPPTTSGPQTHQ
		1				PKEHLTNFKSGVRPRRDACLGP
						SPLAASPAFLGKGQVPQPLISLC
						PDPLFPHPNLISLRPNPLCPHPD
						LVSLCPDPFPAFLEAHKNFQTPE
						PQQPGIPPEPPPGACYKCQKSD
						HQAKECLQPRIPPKPCPICAGPH
		1				WKSDCSTHLAATPRAPGTLAE
		_				GSLTDSFSA
16248	46616	В	16343	1	498	
16249	46617	A	16344	3	903	
16250	46618	A	16345	1	684	
16251	46619	A A	16346	2	1028	
16252	46620	A	16347	1	338	
16253	46621	A	16348	790	392 2184	
16254 16255	46622 46623	A	16349 16350	773	890	
16256	46624	A	16351	467	606	
16257	46625	A	16352	1569	1841	
16258	46626	A	16353	430	980	LLTEDALDDLIPSFLLTGQQTPA
10230	140020	1	10333	130	700	FGRRVSGVIEIADGSRRRKAAA
					İ	LTESNYRVLVGELDDEQMAAL
ļ						SRLGNDYRPTSAYERGQRYASR
						LQNEFAGNISALADAENISRKII
						TRCINTAKLPKSVVALFSHPGE
						LSARSGDALQKAFTDKEELLKQ
1						QASNLHEQKKAGVIFEAEEVIT
						LLTSVLKTSSASRTSLSSRHQFA
		1				PGATVLYKGDKMVLNLDRSR*
						RGIT*AAGI*PS*AEKSWGDI*S*
	L	L	L	L	l	RSYHSFNFCA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
16259	46627	Ā	16354	i i	343	MKRAPVIPKHTLNTQPVEDTSL
10237	10027	`	1055.			STPAAPMVDSLIARVGVMARG
						NAITLPVCGRDVKFTLEV\SGVI
					,	VLRRPLGYGQVMNVTRSCLLR
İ						THWMISSLLFY*LVNRHRRSVE
	}					EYLVS
16260	46628	A	16355	1	584	MLRDLLKNVDLKGFEPDVRILL
1.0200						TKYSNSNGSQSPWMEEQIRDA
						WGSMVLKNVVRETDEVGKGQI
						RMRTVFEQAIDQRSSTDNEGAP
						VIPKHTLNTQPVEDTSLSTPAAP
						MVDSLIARVGVMARGNAITLP
						VCGRDVKFTLEV\SGVIVLRRPL
						GYGQVMNVTRSCLLRTHWMIS
			<u> </u>		į	SLLFY*LVNRHRRSVEEYLVS
16261	46629	A	16356	62	350	SLPNLDNAAICSSSSSPTRTR*SL
						SVSAAALRRRLPSAISDHTREVF
						STLSPRSTSRVNFTSRPHTGKV
						MALPRAITPTRAINESTIGAAGV
						DNEV
16262	46630	Α	16357	291	502	RLFEVHHHVHKHLLGDNRYPI
						WIMQPSAHHP/RSPTRTR*SLSV
1						SAAALRRRLPSAISMTPDTLRP
						NAGVC
16263	46631	Α	16358	1	378	
16264	46632	A	16359	159	433	SYVKYFPHQPAQKYFQQIHSAI
						GLHNADHVHKHLLGDNRYPIW
					}	IMQPSAHHP/RSPTRTR*SLSVS
						AAALRRRLPSAISMTPDTLRPN
						AGVC
16265	46633	В	16360	1	624	
16266	46634	Α	16361	3	299	
16267	46635	Α	16362	163	699	
16268	46636	A	16363	795	980	
16269	46637	Α	16364	663	734	
16270	46638	A	16365	391	1248	
16271	46639	Α	16366	1	1143	
16272	46640	Α	16367	1	271	MOLIDAY MONTO TO THE TOTAL TO T
16273	46641	Α	16368	1	493	MSKRRWLTGNGSDRENIRLDL
						GFLVSDEEKRYKCEDCGKAFN
						RSSNLTTHKKIHTGEKPYKCEE
	1					CGKAFKRSSILTTHKRIHTGEKP
			1			YKCEECGKVFKYLSSLSTHKIIH
						TGEKPYKCEECGKAFNWSSHL
						TTHKRIH\MERNPTNVKNVAKA
	1.55:5	 	1.62.63		1.400	LSTPLPLLNIK
16274	46642	Α	16369	728	1428	
16275	46643	Α	16370	3689	3864	NYKSFFKWS/SQMQSMTKIYRG
						PLDHPDSPCSNVNDIEGTPPEEI
L		<u> </u>	l	<u></u>	L	STAQPLLHPRPAGSS

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	1	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide inscrtion)
16276	46644	A	16371	2	126	WPLLLLGCLSLTKEWGFQA*LE
						KHVKRVKFPSHNLVPGRSI
16277	46645	Α	16372	728	2207	
16278	46646	С	16373	1	873	
16279	46647	Α	16374	1220	2097	
16280	46648	Α	16375	73	1438	QDLRLFYLKDTMFSKLAHLQR
						FAVLSRGVHSSVASATSVATKK
						TVQGPPTSDDIFEREYKYGAHN
						YHPLPVALERGKGIYLWDVEG
						RKYFDFLSSYSAVNQGHCHPKI
						VNAL\KSQVGQN*PLTSRAFYN
ŀ						NVLGEYEEY\ITKLF\NYHKVLP
						MNTGVEAGETA\CKLA\RKWG
						L/YTVKGIQKYK\AKIVFAAGNF
						WGRTLSAISSSTDPTSYDGFGPF
						MPGFDIIPYNDLPALERALQDP
				1		NVAAFMVEPIQGEAGVVVPDP
						GYLMGVRELCTRHQVLFIADEI
						QTGLARTGRWLAVDYENVRPD
						IVLLGKALSGGLYPVSAVLCDD
						DIMLTIKPGEHGSTYGGNPLGC
						RVAIAAL\EV\LEEENLAENADK
						LGIILRNELMKLPSDVVTAVRG
						KGLLNAIVIKETKDWDAWKVC
						LRLRDNGLLAKPTHGDIIRFAPP
						LVIKEDELRESIEIINKTILSF
16281	46649	Α	16376	600	806	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
16282	46650	Α	16377	11	2255	EPDVCAKMAGRSMQAARCPTD
10282	40030		10377	1'	2233	ELSLTNCSVVNEKDFQSGQHVI
						VRTSPNHRYTFTLKTHPSVVPG
						SIAFSLPQRKWAGLSIGQEIEVS
						LYTFDKAKQCIGTMTIEIDFLQK
						KSNDSNPYDTDKMAAEFIQQF
						NNQAYSVGQQLVFSFNEKLFG
						LLVKDIESMDPSILKGEPATGK
						RQKIEVGLVVGNSQVAFEKAE
		1				NSSLNLIGKAKTKENRQSIINPD
1						WNFEKMGIGGLDKG\FSDIFRR
						AFASRVFPPQIVEQMGCKHVK
						GILLYGPPGCGKTLLARQIGKM
			}			LNAREPKVVNGPEILNKYVGES
						EANIRKLFADAEEEQRRLGANS
						GLHIIIFDEIDAICKQRGSMAGS
						TGVHDTVVNQLLSKIDGVEQL
						NNILVIGMTNRPDLIDEALLRPG
						RLEVKMEIGLPDEKGRLQILHIH
						TARMRGHQLLSADVDIKELAV
						ETKNFSGAELEGLVRAAQSTA
						MNRHIKASTKVEVDMEKAESL
		1				QVTRGDFLASLENDIKPAFGTN
						QEDYASYIMNGIIKWGDPVTRV
		1				LDDGELLVQQTKNSDRTPLVSV
						LLEGPPHSGKTALAAKIAEESN
						FPFIKICSPDKMIGFSETAKCQA
		1				MKKIFDDAYKSQLSCVVVDDIE
		İ				RLLDYVPIGPRFSNLVLQALLV
						LLKKAPPQGRKLLIIGTTSRKDV
ľ						LQEMEMLNAFSTTIHVPNIATG
			1.50.70		2101	EQLLEALELLGNLKDKERTTIA
16283	46651	В	16378	1	2181	
16284	46652	A	16379	1	441 482	
16285	46653	В	16380	1	636	MRDPNTKRSRGFGFVTCATVE
16286	46654	Α	16381	I	030	EVDAAMNARPCKVVGRTVEPK
		1				RAVSREDSQRPEDTEELHLRDY
		1				FEQYGKIEVIEIMTDQGSGKKR
						7
						GFAFVTFDNHDSMDKTVIQKY
						HTVNGHNCEARKALSKQEMAR
						ASSSQRGRSGSGNFGGGRGGGF
						GGNDNFGRGGNFSGHGGFGGS
						HGGGGYGG/SGDGYNGFGNDG
1.6267	146655	_	16202	1	760	GGGSYNDFVNYNNQSSHFGPM
16287	46655	A D	16382	50	762	
16288	46656	В	16383	59	1042	
16289	46657	В	16384	76	848	
16290	46658	В	16385	179	400	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
16291	46659	A	16386	27	372	VGSLRRPGPPREETRRLRERRRS LEPWAFLLKEAKVG*QPSSLFQ WEGPMGLCSTPLVMNMGDSLI TK\HLFQVPPLTITSPGPRTRRPH S*YPSPSLATGVWAAVWCKQP HFPAK
16292	46660	В	16387	134	284	
16293	46661	Α	16388	1	2013	
16294	46662	A	16389	313	770	IFGFVVAMHGAPGQAVLELLQ LTGDQSLGLSQRHMAALDALV HLLHGGLAVLTTAGAAPEHET RGIAGLREEQCRNPRCVKRGW AQEKQA*PGQRGGRKSSPFWSF TL/DSASAGPRAAHLHLPDVLS LQTLHHGLHRLLHPELLQLSHG
16295	46663	A	16390	1	2753	
16296	46664	Α	16391	1	1363	
16297	46665	Α	16392	26	181	
16298	46666	Α	16393	1	867	
16299	46667	Α	16394	1	1447	
16300	46668	A	16395	31	659	APALPGCEHMMAIRELKVCLL GDTGVGKSSIVCRFVQDHFDH NISPTIG\ASFMTKTVPCGNELH KFLIWDTAGQERFHSLLPLYL\R GSAAAVIVYDFTEAGFHFHPLK KWV\KRLKELGPE\NIVMAIAG NKCDLSDI\REVPPEGMLKEY\A ESIGAIVVETSAKNAINI\EELFQ GISRQIPPLDPHENGNNG\TIKVE KPTMQASRRCC
16301	46669	Α	16396	1	1182	
16302	46670	Α	16397	1	513	
16303	46671	A	16398	146	429	TFYTCFSPSVIPFSFSIHTKPYPG HHQSFYMTNVSSNNPTISPLTT RPPFSLISPALGSHAAPNPA*SSP EKHRPFSLHTTPQKFHRPNTSTL F
16304	46672	Α	16399	1	435	
16305	46673	A	16400	56	477	ARAAKGESCEGPGVGRCFSFSS YRQTPPCCQRLVNVPKTR\RTF CKK\CGKH\RPHKVTQYKKGKD SFYA\QGRRRYDRKQSGYGGQT KPIFRKKAKTTKK\MGLRLECV EPNCRSKR\MLAIKRCKHFELG GDKKRKGQVIQF
16306	46674	A	16402	183	373	IHGSSENPTPERETLFWHAENG SALASC*KSGGQAPKRKVRTTL ILKAEILILGMLVLRSHGM

16307 46675 A 16403 1 881 MPFKCKKQHTFNKTSPKEGAI FTHGCLCTSPPNSELSEHLGNL KLSSWEHLHRSQQTLSVRLHVPTPOPVEHEFIQLIYHTGAKI QASKVIHATKDGRRRKSPLWQ KYPKLILREASSVSEELHKEVQ EAFLTLHKHGCLFRDLVRIQG DLLTPVSRILIGNPGCTYKYLN RLFTVPWPVKGSNIKHTEAELA AACETFLKLNDYLQETIQALE LAAKEKANEDAVPLCMSADP RVGMGSSYNGQDEVDIKSRA. YNVTLNFMDPQKMPYLKEE YFGMGKMAVSWHHDENLVD IQQQWDPVDGTTASSA VPKQ GRDKRREVPNTLSQPNSQEGL SAFHSRCVHAVFYKLTSAPKT SLPPVPSDLLAEGLAASRAGQ LWRCDKSLLTLVKKINLQGK RDGGRSATNQQSPHLDNTGI ERQLEQKVGRCRSPWRGIGHI EGNAQKGHLLHGVNKNGQHL L*MPSPLPGGGKGRLVGMLS. WVLT*VYKIQHIGT/PAA VEC* EAFLAVWLGEGVGDFSPLVPS LWHS*RGCCSHGIPELELNPV QIFIM/SASSLPFSPCQNK VPLSQ 16309 46677 A 16405 1 2025 16310 46678 A 16406 1 316 16311 46679 A 16407 1 217 16312 46680 A 16408 3 135 135 131 131 131 14679 A 16407 1 217 16312 46680 A 16409 21 218 16314 46681 A 16409 21 218 16314 46682 A 16410 392 3744 LNYKMILLKNLSPMSMLKAFSI AIPMQPPAQTINLPTARPNGFQ LILLSQYNACPPPQ*EAG/QVDL CSTIPLNLLPDSLPLTVPTGPLP GSVGLVLGRASSSAKGITHTGO INSDSVDEIKLITSAKVPVSILA ESIAQLLLLPNILLNRGDKTRGI GMGSRGEKAAYWINVISKQRR TCTIHIQGRFEGLVDTGADVS VISSLWPSSWLKHPTNMGLV	SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
16307 46675 A 16403 1 881 MPFKCKKQHTFNKTSPKEGAI FTHGCLCTSPPNSELSEHLGNN KLSSWEILHHRSQOTLSVRLE VPTPQPVEHEFIQLIYHTGAKI QASKVIIATKDGKRKSPLWQ KYPKLILREASSVSELHKEVK EAFLTLHKHGCLFRDLVRIQG DLTPVSRLIGNPGGTYKYLIN RLFTVPWPVKGSNIKHTEAEIA AACETFLKLNDYLQIETIQALE LAAKEKANEDAVPLCMSADP RVGMGSSYNQDEVDIKSRA-YNVTLLNFMDPQKMPYLKEE YFGMGMAVSWHHDENLVD IQLQLWDPVDGTTASSAVPKQ GRDKRREVPNTLSOPNSQEGI. SAFIISRCVIIAVFYKLTSAPKT SLPDVPSDLLAEGLAASRAGQ LWRCDKSSLTLVKINLQGR RDGGRSATNQSQPHLDKYSMLSVEYKLTSAPKT SLPDVPSDLLAEGLASSRAGQ LWRCDKSSLTLVKINLQGR RDGGRSATNQSQPHLDKYNKNQGHL L*MPSPRLPGGOKGRLVAGHLE EGNAQKGHLHGVNKNGGHL L*MPSPRLPGGOKGRLVAGHL L*MPSPRLPGGOKGRL*MPSPLAGGOKGGGGGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG	NO:	of peptide	hod	1	1	l .	*=Stop codon, /=possible nucleotide
16307 46675 A 16403		sequence		09/540,217		of peptide sequence	deletion, \=possible nucleotide insertion)
FTHGCLCTSPPNSELSEHLGNI KLSSWEHLHHRSQQTLSVRL- VTPQPVPHEPIQLIYHTGAKI QASKVIIATKDGKRRKSPLWQ KYPKLILREASSVSEELHKEVC EAFLTLHKHGCLFREDVRIQG DLLIPVSRILIGNPGCTYKYLN RLFTVPWPVKGSNIKHTEALIL AACETELKINDYUCJETIQALE LAAKEKANEDAVPLCMSADF RVGMGSSYNGQEVDIKSRA- YNVTLINFMDPQKMPYLKEE YFGMGKMAVSWHHDENLVD IQLQLWDPVDGTTASSAVPKQ GRDKRREVPNTLSOPNSQEGL SAFHSRCVHAVFYKLTSAPKT SLPVPSDLLAEGLAASRAGQ LWRCDKSSLITLVRKINLQGR RDGGRSATNQQSQPHLDNTGI ERQLEQKVGRCRSPWRGLGHH EGNAQKGHLLHGVNKNQGHI L*MPSPLPGGQKGRLVGMLS WVLT*VYKIQHI/GT/PAAVEG* EAFLAVWLGEGVGDFSPLVPS LWHS*RGCCSIHGIPELEINPV QIFIM/SASSLPFSPCQNKVPLSG 16308 46676 A 16405 1 2025 16310 46678 A 16406 1 816 16311 46679 A 16407 1 217 16312 46680 A 16408 3 135 16313 46681 A 16409 21 218 16314 46682 A 16409 21 218 16315 46683 A 16410 392 3744 LNYKMLLKNLSPMSMLKAFSI AIPMQPPAQTINLFTARNIGGU CSTIPLNLLPDSLPLTVPTGPLP GSVGLVLGRASSSAKGITHITGI INSDSVDEIKLITSAKVPVSILA ESIAQLLLLPNILINGDKTRGI GMGSRGEKAAYWINVISKQRR TCTHIQGRKFEGLVDTGADVS VISSL WPSSWLKHPTNMGLV VGKAEEVYGSTFILPCTGLDGG VGKAEEVYGSTFILPCTGLDGG VGKAEEVYGSTFILPCTGLDGG VGKAEEVYGSTFILPCTGLDGG VGKAEEVYGSTFILPCTGLDGG VGKAEEVYGSTFILPCTGLDGG VGKAEEVYGSTFILPCTGLDGG VGKAEEVYGSTFILPCTGLDGG VGKAEEVYGSTFILPCTGLDGG	,				sequence		
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16309 46677	16200	16676	_	16404	 	450	QIFIM/SASSLPFSPCQNKVPLSG
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16316 46684 A 16412 1 1911	16315	46683	Α		1	783	
	16316	46684	Α	16412	1	1911	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
16317	46685	A	16413	3	1381	KTPFVGYQHQRPKVDKTTKMG RNQSRKAENSKNQSTSSPPKDR SSLPATEQSWTENDFDELTEVG FRRSAITNFSELKEHVLTRREA KNLEKRLDEWLTRINSVEKTLN DLMELKTMAQELRDARTSFNS *FNQVEKRISVIEDQIDEIKQDD KAAETCNWRWVSESLRQLRAS VDAFHARASHYNAGECLHQLA ALNSRLNCAQEMARRDSIGEVP PVPWRTVVGSGIAGEAKLDHL RLVSLIGREVNKENSPAATRWL FSFQAGALAGGQIVLQAAKPN AHGQPVVATRNLLNPAGIVAV LKQRPRLVAAAVQAFYLRDPID LRACRVFKTFLPETRIMTSYYAI EWSLWTLVDVLHAENSESAHM SHNSYDPPSDVARHLTHLPMSP IPDYIIARFRYRHRSRVRLTARK LSPAFMRCSSEIQRYSTMLERG FNGGYCRSTTVRVVTEATRML
16318	46686	A	16414	1	1191	
16319	46687	Α	16415	I	1909	
16320	46688	В	16416	1	1791	
16321	46689	Α	16417	1	1818	
16322	46690	A	16418	1	1044	
16323	46691	В	16419	1	1140	
16324	46692	Α	16420	1	1062	
16325	46693	В	16421	1	972	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	1	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
16326	46694	A	16422	1	1758	MDTFLQAERKDYMEAYELIEQ
		1				EEQGVKELNRPLTAPSALHPRD
						EAHLIMVDKLFDVLLDSVCQYF
				1		IEDFCIDVHHINSRWI/AVCQYFI
						EDFCIDVHHINSRWIKDLNVRP
						KTIKTLEENLGNTIQDIGMGKD
						FLSKTPKAMATKAKIDKWDLIK
						LNNFCTAKETTIRLPVPNPCGPV
						AETPLSRVFVLREARKTLKPHQ
						STCLSKGRDNSIDSWKTAARVL
						KTAISSTPTIPILKDQTQEWSVS
				:		ATFTSDGKIRLFYTDYSGKHYG
						KQSLTTAQVNVSKSDDTLKING
						VEDHKTIFDGDGKTYQNVQQFI
						DEGNYTSGDNHTLRDPHYVED
						KGHKYLVFEANTGTENGYQGE
					1	ESLFNKAYYGGGTNFFRKESQK
						LQQSAKKRDAELANGALGIIEL
						NNDYTLKKVMKPLITSNTVTDE
				1		IERANVFKMNGKWYLGELGW
						RREWSSGDRKSLARLSKTGTLG
						LAALTYKCTRDQWTVYCRVIR
						EKNLCQDMRWYQRCCQTCRD
						FYANKMSPATAELVTRSPKGRS
						KLRLRYQLVFISLKGKPARKGK
						ASSGSKLSLLSSSKTVALAGMG
						RGDEFQMELESEDLTSERDGIL
						TPTSSNQKKSKAYGSLIRT
16327	46695	A	16423	1	4449	
16328	46696	Α	16424	3	3859	QFLPNLDSTVLGENYFDGTYQ
						MLYLLVKGTIPVEIHTATVIFVS
						FQLSVATEDDFYTSHNLVKNLA
						LFLKIPSDKIRISKIRGKSLRRKR
						SMGFIIEIEIGDPPIQFISNGTTGQ
						MQLSELQEIAGSLGQAVILGNIS
						SILGFNISSMSITNPLPSPSDSGW
						IKVTAQPVERSAFPVHHVAFVS
						SLLVITQPVAAQPGQPFPQQPSV
						KATDSDGNCVSVGITALTLRAI
						LKDSNNNQVNGLSGNTTIPFSS
						CWAN

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
16329	46697	Α	16425	65	1905	RTSSRKTTNHCSEIKEDTNNWK
						NIPCSWIGRINIVKMAKLPKPPL
						VIPRQTGSGVDLQQTPTDLHLR
						VLTVRRKTNKQKGHPHQNPICT
						SPSSKTKVLEVLARAFRQEKEI
						KGIQVGKEEVKLSLFADDMIVY
						LENPIVSAQNLLKLISNFSKVSG
						YKINVQKSQAFLYTNNRQTESQ
						IMTELPFTTASKRIKYLGIQLTR
						DVKDLFKENYKPLLNEIKEDTN
						KWKNIPCSWVGRINIVKMAILP
						KVIYRFNAIPIKLPMTFFIELEKT
			!			TLKFIWNQKRARIAKSILSQKN
						KAGGITLPDFKLYYKATVTKTA
						WYWYQNRDIDQWNRTEPSEIM
						LHIYNYLIFDKPEKNKQWGKDS
						LFNIWCWENWLAICRKLKLDPF
						LTPYTKINSRWIKDLNVRSKTIK
						TLEENLGNTIQDIGMGKDFMSK
						TPKAMATKAKIDKWDLIKLKSF
						CTAKETTISVNRQPTEWEKIFAI
						YSSDKGLISRIYNELKQIYKKKT
						NNPIKKWVKDTNRHFSKEDIYA
						AKRHMTKCSSSLAVREMQIKT
						TMRYHLTPVEEVRAGEMAKTR
		ļ				RCQNVKGASEEGIRA/HCFVSH
						*PHCP/LWRATTLMETLKFKPIE
						VPCKPLEKSVSLLLL
16330	46698	В	16426	1	2348	
16331	46699	Α	16427	1	660	MPNFFIDRPIFAWVIAIIIMLAGG
						LAILKLPVAQYPTIAPPAVTISA
						SYPGADAKTVQDTVTQVIEQN
						MNGIDNLMYMSSNSDSTGTVQ
						ITLTFESG\QVQNKLQLAMP/LL
						PQEVQQQGVSVEKSSSSFLMVV
					}	GVINTDGTMTQEDISDYVAAN
						MKDAISRTSGVGDVQLFGSQY
						AMRIWMNPNELNKVERNSRRQ
						DVGERDISSGSRKVNKESREDE
		<u> </u>	16125			EVT
16332	46700	A	16428	1	975	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	1	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	<u> </u>					
16333	6333 46701	A	16429	1171	2007	AVDIVQGLVRGWGRQGWLLS GRGIPEGPLLLWGATVGSWRR LLPLTGVCGLPLGAVPVLIS*LL GGAQVQLLREEEAGHELVQQL
						VDGALVLLLDGRLPLHVVADE RCCVLQDLPVLVHKDEVRLAR
		İ				GGTAALRVLLLVLVVDALDLV
						EQALGAQGLGVVLLEVHSLVV
		ŀ				KGLEVRLLILLPPDLVEALLGLP
		ŀ				PLLFLGLQSFQDNGDSGDRVLL
			ļ !			EALDDLRGLVAIDAINGTALDH
						LRRASQGVVQPELLLVFRRHPR
						QLLLHTFQHTDCGQVRLQDIV
16334	46702	В	16430	1	768	
16335	46703	A	16431	1	2553	
16336	46704	Α	16432	3	573	CEMHSVQKNEHDKTPEHSRKR
ļ		ł	į			SRPSLLRPVSAPAKKKMKLRGT
ł		1				KDLSIAAVGKYGTL/PGIFFL*Q
						GPPGAEEPGGV*KLPPLHRTLQ
		l				PGAGVWL*APAARQPISGVSS*
		l				LPRAP*GGSVPVNSIYLVFSYCF
			l			*AMIVLF*NPVY*LT*LI*LLIPM
						HL*GAIHHLGTLQLVPLPVTSV VSILTRIQWVLE
16337	46705	A	16433	110	1347	LTTATGMELSMGSTTWTSLCK
10337	40703		10433		1547	AENAACQQEKPHIWLCSISSFL
						RLKRRRTTTKNTGTKKIASTVA
		1				VTIPPITPLPTAFCAPEPAPVLLT
			1	-		SGIPPRLNASENPQLLSGS/SSSD
		ŀ				STGTVQITLTFESGTDADIAQVQ
						VQNKLQLAMPLLPQEVQQQGV
			<u> </u>			SVEKSSSSFLMVVGVINTDGTM
						TQEDISDYVAANMKDAISRTSG
		ŀ				VGDVQLFGSQYAMRIWMNPNE
		İ				LNKFQLTPVDVITAIKAQNAQV
						AAGQLGGTPPVKGQQLNASIIA
						QTRLTSTEEFGKILLKVNQDGS
						RVLLRDVAKIELGGENYDIIAEF
						NGQPASGLGIKLATGANALDT
						AAAIRAELAKMEPFFPSGLKIG
						YPYDTTPVVNISIHEVGKTLVE
		ŀ				ANNLVFLAMNLVLQEFPADVIS
16000	46706	_	16424	1	2550	TIAGPPSLRSSRVPGL
16338 16339	46706 46707	B B	16434 16435	1	2550 630	
16339	46708	A	16436	270	455	
16341	46709	A	16437	174	492	GALQPPPCTVGAPFWAGQGWS
1,0341	10,00	` `	1.0.57	.,,		PLPQLAGRCGGRGT\RGTGAAC
						GACGPAGVPGGLGLGGPRTRSS
						RPALLAPGNGGLSTRASGCGGC
						TESPSSGWPTSAALDFLPGP
	<u> </u>	L	L.,	l		TEGI GOO WI TOAMEDI DI GI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
16342	46710	Α	16438	3	413	
16343	46711	Α	16439	31	507	
16344	46712	Α	16440	1	537	
16345	46713	Α	16441	152	247	
16346	46714	Α	16442	44	208	EEMTSKMYKVIIRKTMKASVC
						TQVKHPAIASICKAVL*VGPWS
			ļ			GHFKYAGLPTL
16347	46715	Α	16443	1	549	
16348	46716	В	16444	376	862	
16349	46717	Α	16445	2	1603	TNVSTTIATVVLNALARATKLL
						PPGHRRSPNPATAAASSPPTTLA
	1					HCSLRSHHQPLRGEQQPQAIRT
						TAGRCKALSSCPSFFECGSRSSG
		į				APPGACGQIFRVSNAKLEPQAR
			ŀ		<u> </u>	PSSPVTGGPLDLMACRVVAAST
						REMAMLIAQALQTINYGRDDE
						NGHTHSACEVHHGGRVPELSV
	İ	İ				PSFSWRNRNNPSFIMGSITPTDY
						TLSKCYLPREDVVLIIYCGVSTN
						EQPDEEVPRTRFPTVAVSIGFAV
		İ				RQELEFGVIYHCLE\YRLYTGRR
						GRGAFCNGQRLRVSGETASGM
						LPGIQQIHIQQEQNLQVGASTEL
						PCNLHTTRIRVASAPSSAPEDGT
		ĺ				ESKTCNFRVTSSKGYCLALPRV
						PFPLTTGYKHLHKPPTGTHSTY
						HSMCLLNQPGWLRTISIILFLNN
						KICWQKKSGREIKIEDYFPEYA
						NYTVPEDGHLSAHLYILSGLCM
						WŁQHSGQDSILYTDYRRGGEN
						SFVMVNSVALNGDGCGICSETE
						AELIEVSHRLNCSREVGEHLNA
						TGELGEERAHGRQQAACGFTIA
16350	46718	Α	16446	1	648	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
,	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
16351	46719	A	16447	329	2497	GEGQAVLQGKMGGEEKPI\GA
						GEEKQKEG\GKKKNKE\GSGDG
						SRAELNPWPEYIYTRLEMYNIL
						KAEHDSILAEKAEKDSKPIKVT
						LPDGKQVDAESWKTTPYQIAC
				ļ.		GISQGLADNTVIAKVNNVVWD
						LDRPLEEDCTLELLKFEDEEAQ
		l				AVYWHSSAHIMGEGMERVYG
				ļ		GCLCYGPPIENGFYYDMYLEEG
		1				GVSSNDFSSLEALCKKIIKEKQA
						FERLEVKKETLLAMFKYNKFK
i		İ				CRILNEKVNTPTTTVYRCGPLID
						LCRGPHVRHTGKIKALKIHKNS
		ŀ				STYWEGKADMETLQRIYGISFP
						DPKMLKEWEKFQEEAKNRDHR
						KIGRDQELYFFHELSPGSCFFLP
						KGVYIYNALIEFIRSEYRKRGFQ
						EVVTPNIFNSRLWMTSGHWQH
						YSENMFSFEVEKELFALKPMNC
ļ		1				PGHSLMFDHRPRSWRELPLRLA
		İ				DFGGLHRNELSGALTGLTRVRR
						FQQDDAHIFCAMEQIEDEIKGC
						LDFLRTVYSVFGFSFKLNLSTRP
						EKFLGDIEVWDQAEKQLENSL
						NEFGEKWELNSGDGAFYGPKI
						DIQIKDAIGRYHQCATIQLDFQL
						PIRFNLTYVSHDGEDKKRPVIV
						HRAILGSVERMIAILTENYGGK
						LAPFWLSPRQVMVVPVGPTCD
						EYAQNVRQQFHDAKFMADIDL
		ļ.				DPGCTLNKKIRNAQLAQYNFIL
						VVGEKEKITGTVNIRTRDNKVH
16352	46720	С	16449	189	456	
16353	46721	A	16450	3	1414	
16354	46722	Α	16451	95	293	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
	of peptide	hod	in USSN	location of first	,	*=Stop codon, /=possible nucleotide
,	sequence		09/540,217		of peptide sequence	deletion, \=possible nucleotide insertion)
,				sequence		
16355	46723	A	16452	<u> </u> 1	1136	MKKRPGHSLFQFQFLMFRLFSE
10333	40723	Λ.	10732	1	1130	ARQPLSEEDVASLSWRPLHLPS
						ADWQRAALSLWTHRTFREVLK
						EEDVHLTYQDWLHLELEIQPEA
						\MLFQILNGRTSTSGRSMSTFSL
						SPRLQGAVTETCRLRVPFLSTH*
						WISTKAQGVMTTQKILIWSLVA
		1				AQEMRILFPDCRRW*LTWSCSK
					:	TS*CLSATPLPRSTSSLRFSADG
						SRL*QAGGAWLPAFRDRGSC*C
						TNGSSSACLRLSSAAAASRQNS
						PSLPDASSSSTWSTLR*ETSAPT
						EVP*HRTSLPTS\QGPPERLSAE
						QRPLPDGR/RSYWPSARRNA/RL
						ILT/FCSGVVAEPGACAALPVEE
						TLPEPAAPGTAEATRRPAVCQR
						VRVGAIGFPFLLLLDGPAVVTS
			:			DLK
16356	46724	A	16453	3003	3482	HRPRTRPCPWRRNHHLGHRAP
10330	40724	^	10433	3003	3462	L*PQRARR*SPNPPEDGSEDEHL
						*QLC
16357	46725	A	16454	598	759	KVILGLWTEEAQVGGQGLRAP
10337	40723	^	10454	1398	133	TPRLLHHVLK*DIKR*AHRPDA
						VAHACNPNTLG
16358	46726	A	16455	1	1767	VAIIACNINIEG
	46727	A	16456	2	572	
	46728	A	16457	1	726	
	46729	A	16458	155	1334	HALGRRGGSQELSAAAGAASP
10501	1072	ļ^ _	10.00		1.22.	SGSERRAPGALPCAWAAALLV
						LGAPPASRPGGFAAGKTMLLK
		1				E\YRI\CMPLTV\DEYKI\GQLYM
						ISKHSH\EQSDRGEGVEVVQKE
		l		1		PFEDPHHGNGQFTEKRVYLNSK
						LPSWARAVVPKIFYVTEKAWN
						YYPYTITEYTCSFLPKF\SIHIET
		1				KYEDNKGSNDTIFDNEAKDVE
						REVCFIDIACDEIPERYYKESED
						PKHFKSEKTGRGQLREGWRDS
				1		HQPIMCSYKLVTVKSEVWGLQ
						TRVEQFVHKVVRDILLIGHRQA
						FAWVDEWYDMTMDEVREFER
						ATQEATNKKIGIFPPAISISSIPLL
						PSSVRSAPSSAPSTPLSTDAPEFL
						SVPKDRPRKKSAPETLTLPDPE
		1				KKATLNLPGMHSSDKPCRPKSE
16362	46730	A	16459	26	355	
110307				1 .	i '	
		Α	16460	86	489	
16362 16363 16364	46731 46732	A B	16460 16461	86	489 201	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
16366	46734	A	16463	58	553	VARSAPPDGAVCAGPGSRRTE
10300	70737	``	10403		333	MAEQSDEAVKYYTLEEIQKHN
		1				HSKSTWLILHHKG\YDLTKFLE
						EHPGGEEVLREQA\GGD\ATENF
) '					EDVGHSTDAREMSKTFIIG\ELH
		İ				PDDRPKLNKPPEP*RRCFKETLI
						TTIDSSSSWWTNWVIPAISAVA
1.60.57	16505	<u> </u>	16161	105	400	VALMYRLYMAED
16367	46735	A	16464	185	489 345	
16368	46736	A	16465			
16369	46737	A	16466	572	1017	
16370	46738	A	16467	911	1013	TOO YOU GETTOOM GWOED GAN
16371	46739	Α	16468	1	157	FGGNQKGERTGQKGKGERGAK
						RKDFPGAVCISALGLLAKGCG*
						SWRPSSSAN
16372	46740	Α	16469	71	369	SCGLSLIKMTTSQKHRDFVAEP
						MGEKPVGSLAGIGEVLG\KKLE
						ERGFDKAYVVLGQFLVLKKDE
						DLF\REWLKDTCG/APNAKQSR
		1				DCF*CLREWCDAFL
16373	46741	Α	16470	1	831	
16374	46742	Α	16471	368	679	SSQLNKAPSLFNPVSEEFCLWL
						LLLQKELLTRITSLEKNTNDLV
						DLKNTA*ELHEAHTSINS*IDQV
		1				EERISECEDHLTEIRHSEKMKTA
						LLK*DMQTRREKKK
16375	46743	Α	16472	536	717	
16376	46744	Α	16473	1	1632	
16377	46745	Α	16474	1	651	
16378	46746	Α	16475	1	903	GGGGRMKLIDYGLSGYQEESA
		1				E\VKAMDFITSTAILPLLFGCLG
						VFGLFRLLQWVRGKAYLRNAV
						VVITGATSGLGKECAKVFYAA
						GAKLVLCGRNGGALEELIRELT
						ASHATKVQTHKPYLVTFDLTDS
						GAIVAAAAEILQCFGYVDILVN
						NAGISYRGTIMDTTVDVDKRV
						METNYFGPVALTKALLPSMIKR
						RQGHIVAISSIQGKM\SIPFRSGI
						CQPSKHATQAF\FDCLAVPEM\E
						QFEIEVTVISPGYIHTNLSVNAIT
1						ADGSRCGVMDTTTAQ\AEALW
						RWPRMFLLLWGRRRKM
16379	46747	Α	16476	359	1094	
16380	46748	Α	16477	1	906	
16381	46749	A	16478	1	371	
16382	46750	Α	16479	11	257	
16383	46751	Α	16480	2	419	
16384	46752	Α	16481	1	2514	

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
16385	46753	Α	16482	262	822	AGCTRKTRIIDVVYNASNNELV
						RTKTLVKNCIVLIDSTPYRQWY
						ESHYALALGRKKGAKLVRVTS
						L*GLWGGQPDSSLLVMKTLSSS
						ATEGREMRAF*AEEGQHWGVQ
						GSRKLPGPAFLP*AHIFVSPFQT
						PEEEEILNKKRSKKIQKKYDER
						KKNAKISSLLEEQFQQGKLLGE
						KADELEVGSRRD
16386	46754	Α	16483	2	799	SQWLKLPSPLPHAGISRY\NWD
						QAP*KTGGPRESPYHKKREVM
			-			KLGRPSCPN/NPKIGPPAGIQHSP
		1				VCRGG*QVNTRAPEVWTVGNF
						\SWGLICRCTP*NKDPSIVVLQC
						HLNNGAWFRTQGPWVEEFCIV
						ASSDQQQPYR\QWVRSPNLCAC
						PLGPAKKGEPKLDSLRKEEIFK
						QKNDLKKLRKEI*LKRKKNCQI
						PAVLPGREQFPARGKLSCRA/SR
	Ì					FQGPGQLWAEQDGFCA*EGQK
	1					SLEFLSFRKNPRPRKGQIKFLVF
	į					VFHPC
16387	46755	A	16484	1	681	
16388	46756	A	16485	287	694	HSSRPAPASAPRRWGLSWSWQ
						RCWC**TPSPSPRPRPAEG\CLR
			İ			GRGRGAGGGLPFRPHGSLVHS
	ł	Ì				GAQPPHFPHLPGGARQTCGAPP
						GSWSCVL/CRPAETGAAAAGSP
						AAAGSTRKRLGRAQGRGCRVA
						ALRLSPTH
16389	46757	A	16486	189	381	QGAGSVGRTGQQLLKPGPFFLE
						KLLSLGCT/LSG*ALRARACCSP
			:			CCRPGAGCQPQAPVAQAGHP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
16390	46758	A	16487	1	2388	QGRLVAKPPRPARSTAADSSTP
		1				RDMLGLRPPLLALVGLLSLGCV
						LSQECTKFKVSSCRECIESGPGC
						TWCQKLNFTGPGDPDSIRCDTR
						PQLLMRGCAADDIMDPTSLAET
						QEDHNGGQKQLSPQKVTLYLR
						PGQAAAFNVTFRRAKGYPIDLY
						YLMDLSYSMLDDLRNVKKLGG
						DLLRALNEITESGRIGFGSFVDK
						TVLPFVNTHPDKLRNPCPNKEK
						ECQPPFAFRHVLKLTNNSNQFQ
						TEVGKQLISGNLDAPEGGLDA
						MMQVAACPEEIGWRNVTRLLV
						FATDDGFHFAGDGKLGAILTPN
						DGRCHLEDNLYKRSNEFDYPS
						VGQLA\HKLAENNIQPIFAVTSR
						MVKTYEKLTEIIPKSAVGELSE
						DSSNVVHLIKNAYNK\LSSRVFL
						DHNALP\DTLKVTYDSFCSNGV
						THRNQ\PRGDC\DGVQINVPITF
						QVKVTATECIQEQSFVIRALGFT
						DIVTVQVLPQCECRCRDQSRDR
						SLCHGKGFLECGICRCDTGYIG
						KNCECQTQGRSSQELEGSCRKD
						NNSIICSGLGDCVCGQCLCHTS
						DVPGKLIYGQYCECDTINCERY
						NGQVCGGPGRGLCFCGKCRCH
						PGFEGSACQCERTTEGCLNPRR
						VECSGRGRCRCNVCECHSGYQ
						LPLCQECPGCPSPCGKYISCAEC
						LKFEKGPFGKNCSAACPGLQLS
						NNPVKGRTCKERDSEGCWVAY
16391	46759	Α	16488	353	1205	IN TRUITERENERS BEST WITH
	46760	A	16489	11	440	FRALTEGDTQLNWNIVSFPVAE
10372	70700	^	10707	11	TTV	ELSHHENLVSFLETVNQPHHQN
						VSVPSNNVHAPYSSDKEHMCT
						VVYFDDCMSIHQCKISC\ESMG
						SHPNIRWVSLMACCECIGPECID
				,		YGSKTVKCMNCMF*RRQMQT
L		L				KAT**NNRYKKLKK

SEQ ID	-		SEQ ID NO:	1	1	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
16393	46761	A	16490	515	1808	KRKQTGEFGFNSFE\WCDSDLC EMNAGMPKGNLNEQDPKHCPE SEKCLLSIEDEESQQSILSSLENH SQQSTQPEMHKYGQLVKVELE ENAEDDKTENQIPQRMTRNKA NTMANQSKQILASCTLLSEKDS ESSSPRGRIRLTEDDDPQIHHPR KRKVSRVPQPVQVSPSLLQAKE KTQQSLAAIVDSLKLDEIQPYSS ERANPYFEYLHIRKKIEEKRKLL CSVIPQAPQYYDEYVTFNGSYL LDGNPLSKICIPTITPPPSLSDPL KELFRQQEVVRMKLRLQHSIER EKLIVSNEQEVLRVHYRAARTL ANQTLPFSACTVLLDAEVYNVP LDSQSDDSKTSVRDRFNARQF MSWLQDVDDKFDKLKTCLLM RQQHEAAALNAVQRLEWQLK
						LQELDPATYKSISIYEIQEFYVPL VDVNDDFELTPI
16394	46762	Α	16491	1	699	
16395	46763	В	16492	6	5954	
16396	46764	Α	16493	22	585	
16397	46765	A	16494	1	837	
16398	46766	Α	16495	1	1422	
16399	46767	Α	16496	13	189	
16400	46768	A	16497	193	366	GRPGIFKSFCRWW*C*FHMEFI WKSSA*/WSGVWHAGNICLSIS NPEDLRASQFCSSG
16401	46769	Α	16498	1	888	
16402	46770	A	16499	3	531	DAWADAWADAWADAWALQS RRRRRRTQLTLSSPHDCYRGLQ SLLLILCKMATLKEKLIAPVAEE \EA\TVPNN\RITVLGDGQVGMA CAISILGKSL\ADELALVDVLED KLKGGNDGSFQHG\SLFL\QTPK \IVGRIKDYSG\TAQF*RIVVGN WQGVRSARRGKVRSNLVQRN VNVFK
16403	46771	Α	16500	2	406	
16404	46772	Α	16501	2	394	
16405	46773	Α	16502	1	399	
16406	46774	A	16503	74	464	FAFNMPEPAKSAPAPKKGSKK AVTKAQKKDGKKRKRSRKESY SVYVYKVLKQVHPDTGISSKA MGIMNSFVNDIFERIAGEASRL AHYNKRSTITSREIQTAVRLLLP G\EL\AKHAVSEGTK\AVTKYHQ
16407	46775	В	16504	84	832	
16408	46776	В	16505	201	829	
16409	46777	Α	16506	1	804	

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
16410	46778	A	16507	3	615	PECHGIDILSSGQNPHIGSLTGR
						VRAIMVGKAKRKPLELPLPRKI
ĺ						LNQKQYRIAGGIEEISATIKDLK
1						DAGVVIPTTSLFDSPIWPVQKT
1	ŀ					DGSWRMTVDYRQINQVATPIA
						AAIPDVASLLKQINTSPDTW/PI
ĺ						RPPISNGD*GVSGR*ACCLEPLA
						GPHR*ITSEASRILEQGPAIFCR*
						LLSF*ETALGLLLGFGGN*TFDY
16411	46779	С	16508	25	153	
16412	46780	Α	16509	1	1317	
16413 46781	Α	16510	1	1557	QVPGCIIGIDILSSWQNPHIGSLT	
						GRVRAIMVGKAKWKPLELPLP
					RKIVNQKQYHILGGTVEISATIK	
		:			DLKDTEAVTPTTSPFNSPIWPV	
						QKTDGSWRMTVDYCKLNQVV
						TPIAAAVPDV/VSLLEQINTSPG
		l				TWFEWSPK\KALQQVQAAVQA
						ALPFGPYDPADPMVLEVSVAD
						RDAIWSLWNAAIGESQRRPLGF
						WSKALLSSADNYSPFERQLLAS
						YWALVETERLTVGHQVTLRPE
						LPIMNWVLSDPSSHKVSGAQQ
						HWKCAVHT/IIKWKWYIRDWA
						QAGLEGTS*LYWPRASRYQQG
			<u> </u>			HQDLFILRSDLPSQVFIRDKLME
						RRNRRTGRTEKARIWEVTDRT
						VRTWIGEAVAAAAADGVTFSV
						PVTPHTFRHSYAMHMLYAGIPL
			 			KVLQSLMGHKSISSTEVYTKVF
						ALDVAARHRVQFAMPESDAVA
		L				MLKQLS
16414	46782	Α	16511	41	428	
16415	46783	A	16512	1	1026	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
16416	46784	I IA	16513	648	1886	LDPEVAWAKWQHSTVKGPQK
10410	40764	A	10313	1040	1000	QFAFSWQGQQYTFTGLPQGYIN
		1				1
		1				SLSLCYNLIPRDLDRFSLPQDIT
						LVHYIDDIMLTGSSEQELAYTL
						DLLVRRLCAKGWEINLTEIQEA
]	STSVKFLRVQWCGACQDIPSK
						MKDKLLHLFPPTTKKKASLFGF
						RRQCIPHLECGPEQEKALQQAQ
						AAVQAAVPLERYDPADPMVL/
						V/ELTWLWPLLSAQFASSGDQH
						*ALHMAPFLGVVSQLPGGKLIIL
						DIFHHGKGRVLFSLE*TLTPDM
						GLPILHIMLLPRLPSVNSQNALS
						TVMPGFTGPGIKGWKWHHSPS
						PLVIH*QNFCFLFP*HYVLLA*R
	İ					S*FQRKEPCHQET*Q*FH*TGS*
						GCQLDTLGSCYF*VNKLRRELQ
						CWLG*LTQTIKMKSVYYSITEN
						CWMKRSPVKRRKILELEEA
16417	46785	A	16514	1	402	
16418	46786	Α	16515	1	396	PTRPGERSVHVEIPRRVGSGAG
		İ				HAQWAGPVRVARPQLNAQLE
						GWLSQVQSTKRPARAIIAPHAG
						YTYCGSCAAHAYKQVDPSIT\Q
						RFRYSYYDESQGEIYRSIEHLDK
						MGMSIIEQLDPVSFSNYLKKYH
16419	46787	Ā	16516	2	358	
16420	46788	Α	16517	2	769	
16421	46789	Α	16518	1	1017	
16422	46790	С	16519	54	294	
16423	46791	A	16520	159	245	GISVSCSMYL*PDRSAGYYGLF
						KDRKEK
16424	46792	A	16521	ı	1227	
16425	46793	A	16522	1	2697	
16426	46794	Α	16523	200	696	YGIVTGPSLCAGDKQPKKQEK
						NPVLVSPEFVDEALCACEEYLS
						NLAHMDID/IGPGGPAVPQPPRA
						GPSSSSVT\PSGPVSPLQLGLWA
						VGRCLGIDPFCLVQPLPLPQASR
						SFTPLFCSGTLEDSHCLRDRVW
				}		ATFLGSWHSIVTKSMDCATKW
						TAHYPKCVLCGC
16427	46795	В	16524	1	960	THE PERSON NAMED OF THE PE
16428	46796	В	16525	1	2183	
16429	46797	A	16526	306	1844	
16430	46798	A	16527	59	189	YNSGCLYGSQCSVCRCL*GVRN
10.50	10,30		10327			QVRSQQQAMMAMAILVNKKG G
16431	46799	A	16528	1	2034	
16432	46800	В	16529	1	855	
16433	46801	A	16530	1	834	
LUTUU	1 10001	1′ 1	1.0550	I.	100 i	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
16434	46802	A	16531	65	172	
16435	46803	Α	16532	321	429	
16436	46804	A	16533	460	647	
16437	46805	С	16534	529	804	
16438	46806	Α	16535	3	433	
16439	46807	A	16536	1	484	
16440	46808	A	16537	511	681	QQGYTPFQGSGRESFLASSSFW WL/PGLPWLAAA*LQALTPSSH SLLSYLCSLLFCHP
16441	46809	A	16538	51	634	PVPTFGLPARHTGAEKRRKNSG GEQLSKNNLYIRGLPPGGTDQD LIKLCQPYGKIVS\TRAFLDKTTI RCKGYGFVDFDSPAAAQKAVA SLKANGVQAQMAKLQERDPA\ GLSVSSLLL\SMYEQELENMLK PFGHVISTRILRDANGVSRGVG FARMESTEKCEVVIQHFNGKYL KTPPGIPAPSEPLLCKIADG
16442	46810	Α	16539	1	759	
16443	46811	Α	16540	74	352	
16444	46813	A	16541	310	747	GSAVYLCYKKSVAKTNTVSYK AGLICRYPQEDYESFSLPESVPL FCLPMGATIECWPSNSKYPLPV FSTFVLTGASAEKVYGAAIQFY EPYSEENLTEKQRLLLGLTSAD GKSDSSKTIHTNKCICLLSHWPF FD\AFRKFLTFLYR\YSISGAHVL PIEKHISHFMHKVPFPSPQRPRIL VQLSPHDNLILSQPVSSPLPLSG GKFSTLLQNLGPENAVTLLVFA VTEHKILIHSLRPSVLTSVTEAL
16445	46813		16542	319	969	
16446	46814	A	16543	131	342	NSDMGLNLGSALTG*PWVAIEF CCFGFFQIIIVFSCLLGWL*VAP HPNLPSSPAEYMDFREAQEGTD VIY
16447	46815	Α	16544	1	1086	
16448	46816	Α	16545	469	759	
16449	46817	Α	16546	3	598	
16450	46818	A	16547	227	604	
16451	46819	Α	16548	416	1814	
16452	46820	Α	16549	763	867	
16453	46821	Α	16550	1	1419	

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
16454	46822	A	16551	915	3125	SLAGMRLTGTVSGGEVRV/TAK AGARVGPPELVGRPLEGECWY FSGVSRTQAQQLLLSPPNEPGA FLIRPSESSLGGYSLSVRAQAKV CHYRVSMAADGSLYLQKGRLF PGLEELLTYYKANWKLIQNPLL QPCMPQKAPRQDVWERPHSEF ALGRKLGEGYFGEVWEGLWL GSLPVAIKVIKSANMKLTDLAK EIQTLKGLRHERLIRLHAVCSG GEPVYIVTELMRKGNLQAFLGT PEGRALRLPPLLGFACQVAEGM SYLEEQRVVHRDLAARNVLVD DGLACKVADFGLARLLKDDIY SPSSSSKIPVKWTAPEAANYRV FSQKSDVWSFGVLLHEVFTYG QCPYEGMTNHETLQQIMRGWF FGCISRSEAVRRLQAEGNATGA FLIRVSEKPSADYVLSVRDTQA VRHYKIWRRAGGRLHLNEAVS FLSLPELVNYHRAQSLSHGLRL AAPCRKHEPEPLPHWDDWERP REEFTLCRKLGSGYFGEVFEGL WKDRVQVAIKVISRDNLLHQQ MLQSEIQAMKKLRHKHILALY AVVSVGDPVYIITELMAKGSLL ELLRDSDEKVLPVSELLDIAWQ VAEGMCYLESQNYIHRDLAAR NILVGENTLCKVGDFGLARLIK EDVYLSHDHNIPYKWTAPEALS RGHYSTKSDVWSFGILLHEMFS RGOVPYPGMSNHEAFLRVDAG
16455	46823	Α	16552	1	990	
16456	46824	Α	16553		2847	MTAREHSPRHGARARAMQRAS TIDVAADMLGLSLAGEPHRACT GLQTTGYPGL\DPSG*TRK*GRG MSALFL*TASAGA*LH*FSSLVA GAPPG*CGNVQVF/WLAHAQA MGTNNPIFLSSIAFFQDSLINQM TQVKLSVYDVKDRSQGTVKSA ESDRVGNITVIGWQMEEKSDQ RPPVTRSVDTVNGRLHPPAPTV MHSLSHRHSKKNSNFRALALM VLPVDESLTEALGIRSKYASLR KDTLLKSVFGGAICR
16457	46825	Α	16554	1	327	
16458	46826	Α	16555	259	3109	
16459	46827	Α	16556	262	3159	
16460	46828	A	16557	·	1398	
16461	46829	Α	16558	123	393	
16462	46830	Α	16559	245	424	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
16463	46831	Α	16560	1	531	
16464	46832	Α	16561	2348	2775	
16465	46833	A	16562	373	555	LRIQVTFDDVAVEFTPEEWALL DTTQKYLYRDVML\ENYMNLA SVGKSNITLLGMVACTCS
16466	46834	Α	16563	165	457	
16467	46835	A	16564	310	349	K*TMK*EMGKFK*FY/IYLEQED YANANALLSPAPALPKEKRTDS NGRVYFVNHNTRITQWEDPRS QGQLNEKPLPEGWEMRFTVDG IPYFVDHNRRTTTYIDPWSTRP
16468	46836	Α	16565	1	2217	
16469	46837	Α	16566	1402	1860	
16470	46838	A	16567	461	696	KRMCFCTSLSSARSSGSMYMSC RICCSMMPPFSSVFRSISSRYSR AAFW\AWTETRLCWQAAMPRQ QPPGLLHQWTTSP
16471	46839	Α	16568	1	601	
16472	46840	Α	16569	468	1213	
16473	46841	A	16570	2	365	EGCFQKIKLDHILSPPPMPFWK CSNPDVAFGPGKSLKYKRQLS* DGRQLRRGSLGGALTGRYLLP NPVAGQAWPASAETSNLVGMR SQALGQSAPSLTASLPCVSKCW ENIPPWMNCPPIK
16474	46842	С	16571	220	408	
16475	46843	A	16572	840	1091	QSLQHDVTL*VPVLLSYPEPAPS GHGLLRGSAQIAL*TIPADPAGT YLKTREPTLGCQTVPLPAPHLS VDTVCRCLSSSVWPQ
16476	46844	Α	16573	1	1356	
16477	46845	В	16574	1	1084	
16478	46846	A	16575	2	526	QPRGKKFSKNKPRKQNFPTPKH MRVVLPPF/PPLKFFIFPKGLKFL EGGCPNISPPQKKGSLPKIPR*V LIRPPIRKKLFPCTARVKLGPPQ GSFKTAAPSSSSP*SADPVYLAG K/PRQSSSPGGTGGLNPPPRRLQ PQPHPGPSWHSSSSRGPGGHPR RESAERSRCASGNPGGQ
16479	46847	A	16576	36	167	RQLFENTQSDETKQEQNKQNK K*STATESRKVPQTGKSKSYWP
16480	46848	Α	16577	171	527	
16481	46849	В	16578	1	471	
16482	46850	Α	16579	1	294	
16483	46851	A	16580	185	299	GLQYLKGKSGLKVKRGKVW*S ACCKKKGAALGTVASPL
16484	46852	Α	16581	3	414	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	1	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
16485	46853	Α	16582	462	928	WGVPGLPGRHLLPLPSQPPPSW
	1.0000					LSILYVSWAALTCFLNSGESTA
						AKFFCPNLSAISYHTEAIPTTWQ
		1				GVHLSWAFGNGAPDIFSGPGGL
	1					L*PA/PQP/DLALGALFGAGVLV
	-					TTVVAGGITTLLLFIQHHTTQEI
		1				H*QIVQWIKTTL*ILATVSYVIG
16486	46854	A	16583	1	654	
16487	46855	A	16584	66	323	RVSLEIHVIQKRRPHHLSLLSFW
10.07	1.0020	-				TRMKTFRRRK/WGEAEESATDT
						TSETNKVGTGVFVLSLWVPVL
						VRNLAGPGVVLGNICKHFPVC
16488	46856	A	16585	1	1530	
16489	46857	A	16586	615	873	FVFLWSLMMVMYRWGFGVAV
10 103	1.0007	`	1.0000			LSVC*FSF*QSGPSAAGLLEFAG
ĺ						GPFQTPFAWVSVVEAAEQRIML
	1					NSKRCCVIVPLVVSSQRGTRP
16490	46858	Α	16587	286	626	PCKLFPCCNTCYS\RV*DFFRVA
10170	10050	`	10507			GKRSL\LNNYTSWVNTSMLVPR
						VAFACGFPLQQAAVLCVHLLG
						SPILGAADSSVTPPFCWMANAH
						ICGLQLQNTLSFAEPIYLGQEHA
						QLS
16491	46859	A	16588	226	489	PYLSVIFFFDYGINEEQCQRPIQ
10.71	1.0057		1.0500			GDGQNFHRHNGWIKINPSCISF
1						AIFFPFGAEKNMWSTPVTEAFN
						LLST*AF*LFGYKCTLVISSS
16492	46860	A	16589	1	1344	
16493	46861	A	16590	1	257	
16494	46862	Α	16591	73	374	ESETEHKRPGKHSPKRSCNMCG
						QWAFHASLTVSYCLLTWCNET
						CSGDGCEGGMRIDIDPPVSQTR
						FFITLL*LC\FPGAGPQPAYFCTQ
						TQCHVETPRVSQ
16495	46863	A	16592	1	666	
16496	46864	A	16593	95	1840	
16497	46865	Α	16594	1740	2026	
16498	46866	Α	16595	39	487	SRLDPRVRLLFRQFCETRPGLEC
						YIQFLDSVAEYEVTPDEKLGEK
						GKEIMTKYLTPKSPVFIAQVGQ
						DLVSQTEEKLLQKPCKELFSAC
						AQSVHEYLRGEPFHEYLDSMFF
				1		DRFLQWKWLE/RVCACQVRAT
						GK\MYACSRLEKKRIQKRK
16499	46867	A	16596	1	591	
10799	1,000,	1, ,	1.0000	<u> </u>	1	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
16500	46868	A	16597	71	1994	EQEQRGRQGGLLAPPAPAAAA
						AARAAAAAAAAPQALTAPPAG
						SVADRRLSMELENIVANTVLLK
						AREGGGGKRKGKSKKWKEILK
						FPHISQCE\SLRRTIDRDYCRVC
						DKQPIG\RLLFRQFCETR\PGLEC
						YIQFLDSVAEYEVTPDEKLGEK
						GKEIMTKYLTPKSPVFIAQVGQ
]			DLVSQTEEKLLQKPCKELFSAC
				Ì		AQSVHEYLRGEPFHEYLDSMFF
						DRFLQWKWLERQPVTKNTFRQ
						YRVLGKGGFGEVCACQVRATG
]					KMYACKRLEKKRIKKRKGESM
						ALNEKQILEKVNSQFVVNLAY
						AYETKDALCLVLTIMNGGDLK
						FHIYNMGNPGFEEERALFYAAE
						ILCGLEDLHRENTVYRDLKPEN
						ILLDDYGHIRISDLGLAVKIPEG
		1				DLIRGRVGTVGYMAPEVLNNQ
						RYGLSPDYWGLGCLIYEMIEGQ
						SPFRGRKEKVKREEVDRRVLET
						EEVYSHKFSEEAKSICKMLLTK
						DAKQRLGCQEEGAAEVKRHPF
						FRNMNFKRLEAGMLDPPFVPD
						PRAVYCKDVLDIEQFSTVKGVN
						LDHTDDDFYSKFSTGSVSIPWQ
E						NEMIETECFKELNVFGPNGTLP
						PDLNRNHPPEPPKKGLLQRLFK
						RQHQNNSKSSPSSKTSFNHHINS
16501	46869	Α	16598	1	963	
16502	46870	Α	16599	192	609	RPKGARELRLHPGPRYGRTSPQ
i						NCPWARPGCLHEAL*G\WKICL
						HCKCPQEEHMVTVMPLEMEKT
						ISKLMFDFQRNSTSDDDSGCAL
						EEYAWVPPGLKPEQVHQYYSC
						LPEEKVPYVNSPGEKLRIKQLL
	1.60-6	<u> </u>	1.5500		0.5	HQLPPHDNEV
16503	46871	A	16600	30	915 820	
16504 16505	46872 46873	B A	16601 16602	99	248	RLEKHINCSRVI*SWAASPGLTA
16303	40873	A	10002	199	240	AAYRVTLNPPGTFLEGVAKVG
						QYTFT
16506	46874	A	16603	1	717	QTIFT
16507	46875	A	16604	271	556	TAACLQSHHFKSSGQLSFERKW
10307	100/3	~	10007	-''		PKVGTITRSIASLLSGPVFGLHH
						LAFSAHVPREGPTTPWNYFPRV
						AAP/RRWTPGEHARTTTGLGAA
						PCVLLWA
16508	46876	С	16605	135	341	I C V EL W A
16509	46877	A	16606	125	226	
16510	46878	A	16607	194	547	
10210	1-00/0	1~_	110007	134	J71	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
16511	46879	A	16608	2073	2338	
16512	46880	Α	16609	1127	1282	
16513	46881	С	16610	1	2094	
16514	46882	A	16611	523	706	QHPLPNPCCHQPFHLCLRR*TL HCLRQQ*WPPLRQLPGKIMLIL LRSHPQHPCLLLDLQLD
16515	46883	В	16612	190	805	
16516	46884	В	16613	90	2812	
16517	46885	A	16614	44	216	RCRRRLIPHPRRLSPVVIRRIQVP Q/VQPGAGPSRPGHTRDPIGAA AAHCPCPGDSGAH
16518	46886	A	16615	1	1107	MTEEPLQALVLSAVLLYQEAY VRRSLTALGLDTFANFAHFSEM QPLSDAPCGLVTGVTRPLVAYS VEVGGFTIICGLSLASGNISGWC QLYLLLTQCRRGAATLWDSGM GGGRQSSRQTSVPRWTLDSKCI PGPKLTTRPQMDTKALGGQQA HVEHQTHEDIRPQVDTQAPGG HKALGEQQDPVGHQASVGYSP QVPQDSLGQALAGQATPEIPLG LQLHTVLVQEIQELIEAQTRAP GPCAEVRALPAPAAEPEPAWEE APPERALELEGAPAKDQTNEEL PEITA/PYC/EPLALTLELKAWLE RKVGGRG\DQHSPSQQLPCCP*S WARWQTCRQRAGHLAWPPVP RCREASLIH*NHSPAAAGPFILL
16519	46887	A	16616	1	792	
16520	46888	Α	16617	1	963	
16521	46889	Α	16618	1	378	
16522	46890	A	16619		494	MKPRTLAVSVTVLKGGVSRVC SFWCSDVFRVSSFWWVRGLAD SGVKLRTFAVSVTPLKAARLEL FVPPGGFVVSLASGVKLQPFEV SVTAHKGSVDPKSEKQQDLLQ RGKEQSYHTGEGHPSRLP/PAG SGSLLLFSYLAPPTSC*LVQPSG LF*QGADWCIYNP
16523	46891	A	16620	137	262	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
16524	46000	<u> </u>	1.6621	1	2797	EVEL BEATORY BREEF BOY
16524	46892	A	16621	1	2786	EVSLREATQRKLRRFSELRGKL
						VARGEFWDIVAITAADEKQELA
			ļ			YNQQLSEKLKRKELPLGVQYH
						VFVDPAGAKIGNGGSTLCALQC
1						LEKLYGDKWNSFTILLIHSDEW
		ĺ				KKKVSESYVITIERLEDDLPIKE
						KELTELRNIFGSDEAFSKVNLN
						YRTENGLSLLHLCCICGGKKSH
						IRTLMLKG\LRPSRLTRNGFTAL
						HLAVYKDNAELITSLLHSGADI
		ŀ				QQVGYGGLTALHIATIAGHLEA
16525	46000	ļ. —	16622	174	722	ADVLLQHGANVN
	46893	A	16622	174	722 1554	
	46894	A	16623 16624	1	843	
16527	46895	A		1	745	ECWDGEHDIETPYGLLHVVIRG
16528	46896	A	16625	2	143	
						SPKGNRPAILTYHDVGLNHKLC
		1				FNTFFNFEDMQEITKHFVVCHV
						DAPGQQVGASQFPQGYQFPSM
						EQLAAMLPSVVQHFGFK\YVIGI
						GVGAGAYVLAKFALIFPDLVEG
						LVLVNIDPNGKGWIDWAATKL
						SGLTSTLPDTVLSHLFSQEELVN
						NTELVQSYRQQIGNVVNQANL
						QLFWNMYNSRRDLDINRPGTV
						PNAKTLRCPVMLVVGDNAPAE DGVVECNSK
16529	46897	A	16626	3	949	DOVVECNSK
16530	46898	A	16627	1	633	
16531	46899	Α	16628	2	1324	
1	46900	Α	16631	1	591	
16533	46901	Α	16632	138	832	
16534	46902	Α	16633	55	586	IFVHPTPAASTMPSKFDPNEIKV
						VYLRCTGGEVGATSALAPKIGP
			,			LGLSPKKVGDDIAKATGDWKG
						LRITVKLTIQNRQAQIEVVPSAS
						ALIIKALK\EPPRDRKKQKNIKH
						SGNITFDEIVNIARQMRHRSLAR
						E\LSGTIKEILGTAQSVGCNVDG
		L				RHPHDIIDDINSGAVECPAS
16535	46903	Α	16634	1	735	
16536	46904	Α	16636	3	346	
16537	46905	Α	16637	57	649	GPRRAYGGRMAGGGGDLSTRR
						LNECISPVANEMNHLPAHSHDL
						QRMFTEDQGVDDRLLYDIVFK
						HFQR\NKVEISNAIKKTFPFLEG
						LRDRDLITNKMFEDSQDSCRNL
						VPVQRVVYNVLSELEKTFNLPV
						LEALFSDVNMQEYPDLIHIYKG
	I	ł	ŀ	1	1	CENTRAL DI OECCEREDEEDO
				ļ		FENVIHDKLPLQESEEKEREERS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence		*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
16538	46906	A	16638	1	1665	
16539	46907	Α	16639	348	1534	
16540	46908	Α	16640	1	627	
16541	46909	Α	16641	1	1343	
16542	46910	A	16642	76	479	IRAVSRSRERPGSRSRYLQAC*C GGAFPGP*SAEISRLVRSHNLGC YSCT/EGGGGCRKGSCLLQHSC LLWHSCLLPASKNTVCSKDLG GCSCTQESRAPACSWSPRTQGC SASSHDLGSCRNAWETPSRQLC RSW
16543	46911	A	16643	130	675	
16544	46912	C	16644	118	420	
16545	46913	Α	16645	1	987	
16546	46914	Α	16646	1	494	
16547	46915	Α	16647	360	1510	
16548	46916	A	16648	375	2187	VTAAAAATAMAESGESGGPPG SQDSAAGAEGAGAPAAAASAE PKIMKVTVKTPKEKEEFAVPEN SSVQQFKEEISKRFKSHTDQLV LIFA\GKILKDQDTLSQHGIHDG LTVHLVIKTQNRPQDHSAQQTN TAGSNVTTSSTPNSNSTSGSATS NPFGLGGLGGLAGLSSLGLNTT NFSELQSQMQRQLLSNPEMMV QIMENPFVQSMLSNPDLMRQLI MANPQMQQLIQRNPEISHMLN NPDIMRQTLELARNPAMMQEM MRNQDRALSNLESIPGGYNALR RMYTDIQEPMLSAAQEQFGGN PFASLVSNTSSGEGSQPSRTENR DPLPNPWAPQTSQSSSASSGTA STVGGTTGSTASGTSGQSTTAP NLVPGVGASMFNTPGMQSLLQ QITENPQLMQNMLSAPYMRSM MQSLSQNPDLAAQMMLNNPLF AGNPQLQEQMRQQLPTFLQQM QNPDTLSAMSNPRAMQALLQI QQGLQTLATEAPGLIPGFTPGL GALGSTGGSSGTNGSNATPSEN TSPTAGTTEPGHQQFIQQMLQA LAGVNPQLQNPEVRFQQLEQ LSAMGFLNREANLQALIATGG DINAAIERLLGSQHHSSISVS
16549	46917	A	16649	269	440	
16550	46918	A	16650	469	746	RLNLPGPLCAGTSILFAVLMEAI GVSCIRQAGGGHVASHIY*GLM /PALAL*DTEPRNPSALPETCLC GSGPQSQMGHQTSHLLSSPGTG LSF
16551	46919	Α	16651	1	1026	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
16552	46920	Α	16652	719	906	
16553	46921	В	16653	1	1971	
16554	46922	Α	16654	200	239	
16555	46923	Α	16655	109	177	
16556	46924	Α	16656	1	1542	
16557	46925	Α	16657	1	453	
16558	46926	Α	16658	2	487	
16559	46927	Α	16659	1	1260	MEDAEGTVTLLAVTLVARDLN
					i	CIQPQTGPSGGIPEESIVIIGDVSS
						MHVIAPEDLPVGQDVEVEDSDI
		1				DDPDPVRALPLQIHACGRVGFE
						AHDDWSEGGDPAADTLIDPSQ
		1				DARYLKCEGMQAPHNSNPVAQ
		1				KTEISYHIWIARIPREKSGKVTG
						TLCDLLKDLTVIHSLIPKQGCSD
		1				GLGVLVQGTFQPQGNTDTCTC
						QVSAVEGKGEGLMQLASLSRR
						VTIWAYGDSQERDLKTQLEKIT
		1				AIKDMKEMQEKEYRIMEKAHS
						RORKOHHKGPERKEQAAIFAV
	1					LQPPLVITRQTGSGVDPQKTPA
						DLQKRCRTVRRKTSQQKAVAS
						TLTKRITTQKLNAKDHNSSPTR
Ì						EQNCTENEFDELTEVGFRRWVI
						TNSSELKEHVLTQCKEAKNFDK
						RLEELLIQITSLEKNISDLMEL*S
						FALSFCVVILFVNVDATAFCWL
						VFLLTVRHLFCRSAGVFWGSTP
						DPVCLVITSGGCRTAKIAACSFL
						SGPLWCCFLCLEWAFSIILYSFS
						CISFMSLIAVIFSNWVLRSLS
16560	46928	A	16660	1	799	
16561	46929	A	16661	1	969	MHVIAPEDLPVGQDVEVEDNDI
						DDPGQAKKTMEEQWTLVSLLS
		1				TSQQQGPVPPGSTASSATDKLS
						RWELTLLYSTTLFLAREAFRRA
		1				CLSGGTQRDWSQTLNLLWLTV
						PLGVFWSLFLGWIWLQLLEVPD
		1				PNVVPHYATGVVLFGLSAVVE
		1				LLGEPFWVLAQAHMFVKLKVI
						AESLSVILKSVLTAFLVLWLPH
						WGLYIFSLAQLFYTTVLGLCYV
						IYFTKLLGSPESTKLQTLPVSRIT
						DLLPNITRNGAFINWEEAKLAW
						GFFKQFFLKQI\LTEGGRYVGG
						HFLNVLNFGDQGVYDIMNNLG SLVARLIFQPIEESFIYS
16560	146020	_	16662	,	1665	SEVARLIFYFILESFITS
16562	46930	A	16662		1665	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
16563	46931	A	16663	l	1521	MDVSSPPCVDGVSILPPLLIPRQ
		ŀ	-			TGSGVDLQQNPTDLQLRVLTV
		l				KRKTNKQKGHPHQNSICTSPSS
İ		l				KTKEYTFFSVQHHTYSKIDHIV
						GSKAVLSKRKTIEIITNCLPDHS
				ŀ		AIKLELRIKKLTQNHKTTWKQN
						NLLLNDYWVHNKMKAEIKMFF
						ETNKNKDTTYKNLWDTVKAV
						CRGKFIALNAHKRKQERSKIDT
						LTSQLKELEKQEQTHSKASRRQ
		l				EITKIRAELKEIETQKTVQKIHES
						RSWFFEKINKIDRPLAGLIKKKR
		l				EKSQKDAIKNDKLDITTDPTET
				}	-	QTTIRDYHKHLYAIKLENLEEM
						DKFLDTYTLPRLNQESLNRPITG
						SEIEAIINSLPTKKKYRTRRIHSR
		ŀ)
						ILSEVQGGAVLEVLARAIRQEK
		ŀ				EIKCIQLGKEEVKLSLFADDMIV
1						YLENPTVSAQNLLKLISNFSKVS
1					:	RYKINVQKSQAFLYTNNRQTES
						QIMSELPFTIASKRIKYLGIQLTR
						DVKDLFKENYKPLLNEIKED/N
		<u> </u>				KWKSIPCPRIGRINIM
16564	46932	A	16664	2	2315	
16565	46933	A	16665	1	912	
16566	46934	A	16666	1	912	
16567	46935	В	16667	199	619	
16568	46936	В	16668	64	1815	
16569	46937	A	16669	1	239	TRED OVER A COMPRODUM LOVER
16570	46938	Α	16670	1	242	FREDGKTACGHEPSPVLLGLRL
						GLQPWG*GWGCSHGAGSAPDL
						VAGSAHSCGLWLPGSLHHHPA
İ		ļ				QCLPALLCGHLCLSVQDQQNG
1.55	1.505	<u> </u>	1.6691			LLGR
16571	46939	Α	16671	58	505	ASSRTRPSCTLARGRRWFPQPL
						SGRQRQRGGRRGSRVG/YGTA
						QNSTTLTSPAASASEATVSGHW
			}			PSTAPCCV\PAGAG\SATMWRSP
1						A*AACGSAAWTSTWSRRSPGP
		1				GAAAGYGGVRTFCQPAGLTR/L
						ANRSTAGLCGWGLTLSCRETEL
16572	46940	A	16672	242	2952	
16573	46941	В	16673	474	2690	
16574	46942	A	16674	97	451	
16575	46943	A	16675	667	893	
10373	J-0243		1100/3		073	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
16576	46944	A	16676	3	639	RIRVHPTPAAFTMPPKFDPNEIK
10370	40744	^	10070	[037	VGYLRCTGGEVGATSALAPKIG
						PLGLSPKKVGDDIAKATGDWK
						GLRITVKLTIHNRQAQIEVAPSA
						SALIIKALKEPPRDRKKLKILKL
						SGNITFDENGNICSTDMR/HRSL
						SRELSGTTKEILGTAQSVGCNV
		1				DGRHPHDIIDDINSENDKQPRKS
						KMRTSFSPQIPGHEHFHYLLLL
						QHVHCQLALTL
16577	46945	A	16677	350	984	VYLEVAPGGESRLPLSALGPQR
10377	40943	Α	10077	330	704	IGPPGSCLPKKSWVNDICQRQP
		İ				G\DWKGP*GLQVKLTHSRTGQ
						APDCTRVP/VLPSCP*FIQSPSRE
:		1				PPKRPEKKPEKTFKHQLGIFTFL
						NEICSTLLRQMR\HRSLSPEKLS\
						GTPLKEIPGGLPQSSWAGNV*W
						AAIPHDI\IDDIQQWVLWKCPAV
		İ				SDIFIVTVGVKGGPPSVFTEISW
		1]		1
1,6570	46046		16670	262	064	EVLEMVTGGVG
16578 16579	46946 46947	A A	16678 16679	252	964 182	
16580	46948	A	16680	275	639	
16581	46949	A	16681	376	418	
		-	16682	1135	1591	
16582 16583	46950 46951	A A	16683	3	420	AAKEIEVGGGRKAI\KSFQKIQV
10383	40931	^	10005	3	420	RLVRELEKKFSGKHVVFIAQRR
						ILPKPTRKSRTKNKQKRPRSRTL
						TAVHDAILEDLVFPSEIVGKRIR
						VKLDGSRLIKVHLDKAQQNNV
		1				EHKVETFSGVYKKLTGKDVNF
						EFPEFQL
16584	46952	A	16684	1	879	D. I. D. Q.D.
16585	46953	A	16685	2	691	HEEKAMFSSSAKIVKPNGEKP\
10000	'0755	l'`		[~		DEFESGHLPGLFLELE\MNSD\L
				j		KASAQGT*IITAS*RKLKVGGCS
						ESLFH*SFVPRSLKLKTFPQKNP
						KVRA*LRRIGKKKVPVGKAMS
						VLYPPSEGEFLPLSPTSEKARTK
						K*AKSRPPGARTLDKLVH\DAH
						PLRDFGPFPSGNFWAKRIPRSNL
						DGQPGSIKVPFWTKAQQNQCW
						NTRVETFFLVSYK\KLTGQGC*F
						LEFPRVFNC
L		L	<u> </u>			LETTRYTING

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	T .	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
16586	46954	Α	16686	1	452	TLLVIPRKMRSAVELQQTSTDL
						QLRDLTVRKKTNKRKATASTS
						QKGHLHQNPICRSPTSKTKEVG
						FRRSVITNFSELKEDVRTYHKE
						AKNLENRLDEWLTRINSVEKTL
						NDLKELKTMARDLHDACTSFN
	1					RRFDQVEERVTVTEDQIN*QTE
						SNSINFTKRSSTPKPHL\RSPTSK
						TKEVGFRRSVITNFSELKEDVR
						TYHKEAKNLENRLDEWLTRINS
		ł				VEKTLNDLKELKTMARDLHDA
		}				CTSFNRRFDQVEERVTVTEDQI
16587	46955	Α	16687	1	618	
16588	46956	Α	16688	280	353	SCNSIFARSVRACCS*VASSSFC
						C
16589	46957	Α	16689	1	459	IKYSSPMKSTFVDPVDPVAVAV
						IYCLAAGLVIIRIPFFQLVPLKEE
						KMVLLNMDYYEMASLSLVTH
		1				RNEAPFSGRVRSSLPATEQSWM
						ENDFDKLTEVGFRRSVITNFSEL
						KEHVLTHHKEAKNLEKRLDKW
						QTRINSVEKTLNDLMELKTMA*
						EEKMVLLNMDYYEMASLSLVT
						HRNEAPFSGRVRSSLPATEQSW
		l				MENDFDKLTEVGFRRSVITNFS
1						ELKEHVLTHHKEAKNLEKRLD
1	Ì					KWQTRINSVEKTLNDLMELKT
16590	46958	Α	16690	2	403	LALSGEMTLVLFMLLAGGWVT
						HDKPEPMPRAATSGGTRKGKT
	ĺ	1				KAALKDLIATWKEVCVSSPATE
						QSWMENDFDELTEVGFRRSVIT
						NFFKLKEYVLTHHKEAKNLEK
						MLDEWLTRINSVEKTLNDLME
						LKNMA*ELTEVGFRRSVITNFF
	İ					KLKEYVLTHHKEAKNLEKMLD
		ļ.,.				EWLTRINSVEKTLNDLMELKN
16591	46959	A	16691	1	676	
16592	46960	Α	16692	1	1965	
16593	46961	C	16693	1	3075	CALL ORDER DATE CONTROL OF
16594	46962	A	16694	78	152	CAAGFGLPVFY*GFLHRCSLGIL V
16595	46963	В	16695	1	904	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				Sequence		
16596	46964	Α	16696	1	2026	EQEASYGRDAFAKAIYERLFC
						WIVTRINDIIEVKNYDTTIHGKN
	:					TVIGVLDIYGFEIFDNNSFEQFCI
						NYCNEKLQQLFIQLVLKQEQEE
						YQREGIPWKHIDYFNNQIIVDL
						VEQQHKGIIAILDDACMNVGK
						VTDEMFLEALNSKLGKHAHFSS
						RKLCASDKILEFDRDFRIRHYA
	ı					GDVVYSVIGFIDKNKDTLFQDF
		1				KRLMYNSSNPVLKNMWPEGKL
						SITEVTKRPLTAATLFKNSMIAL
						VDNLASKEPYYVRCIKPNDKKS
						PQIFDDERCRHQVEYLGLLENV
						RVRRAGFAFRQTYEKFLHRYK
						MISEFTWPNHDLPSDKEAVKKL
						IERCGFQDDVAYGKTKIFIRTPR
					:	TLFTLEELRAQMLIRIVLFLQKV
						WRGTLARMRYKRTKAALTIIR
						YYRRYKVKSYIHEVARRFHGV
		1				KTMRDYGKHVKWPSPPKVLRR
						FEEALQTIFNRWRASQLIKSIPA
						SDLPQVRAKVAAVEMLKGQRA
						DLGLQRAWEGNYLASKPDTPQ
						TSGTFVPVANELKRKDKYMNV
						LFSCHVRKVNRFSKVEDRAIFV
						TDRHLYKMDPTKQYKVMKTIP
		1				LLQLTCLN\VSNGKDQLVVFHT
	1					KDNKDLIVCLFSKQPTHESRIGE
						LVG\VLVNHFKSEKRHLQV\NV
						TNPVQCSLHG\KKCTVSVETRL
		1				NQPQPDFTKNRSGFILSVPGN
16597	46965	A	16697	1	1665	
16598	46966	Α	16698	1727	1908	FLFFYIC*RELYFQLCGQFWNK
		1				CNVLLRRMYILLIWGGEFCRYQ
						LGLLGAELNSVPGYHC
16599	46967	Α	16699	1	384	
16600	46968	Α	16700	1	636	
16601	46969	Α	16701	1744	2283	
16602	46970	A	16702	194	368	
16603	46971	C	16703	62	205	
16604	46972	В	16704	1	1257	
16605	46973	Α	16705	1	1077	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence	:	
16606	46974	A	16706	1213	1228	AQN*QTQRTSTPKPHL\SSPSTK
10000	46974	A	10/00	1213	1228	
		ŀ				TKENDFDELREEGFRRSVITNFS
	1	ŀ				KLKEDVRTHRKEAKNLEKRLD
	ľ					EWLTRINSVEKTLNDLMELKT
	Ì	İ				MAQELRDTYTSFSSRFDQVEER
	Ĭ	ľ				VSVIEDQMNDMKREEKFREL/K
]				RVQRNEQSLQEIWDYVKRTNL
						RLIGVPEKIQTTIREYYKHLYAN
						KLENLEEMDKFLDTYTCPRLN
		l				QEEVESLNRPVTGSEIEAIINSLP
		1				TKKVQDQADSQPNSTRVFKTV
	1					CISSLGHTTASADTQDVCQLPV
		1				TSLLEQGLSSKSAGFLDYAVQR
						LWLLGTEPL
16607	46975	Α	16707	33	263	
16608	46976	Α	16708	204	1756	
16609	46977	Α	16709	760	821	TMRTHGHRKGNI*QR/R**PEST
						KNSNKFTRKNQTTPSKREWMK
						LETIILSKLSQGQKSKHRMFSLI
						SGN*TMRTHGHRKGNITHWGL
						LWG
16610	46978	A	16710	1	879	
16611	46979	Ā	16711	311	514	KNAHHQWPSKKCKSKLQ*DTIS
		ł				HQLEWQSLKSQETTGTRMKME
						TITLSKLSQGQKSKHCMFSLIGG
						N
16612	46980	A	16712	24	1268	AGLCAARARMRLFRWLLKQPV
						PKQIERYSRFSPSPLSIKQFLDFG
						RDNACEKTSYMFLRKELPVRL
	}					ANTMREVNLLPDNLLNRPSVG
	1					LVQSWDMQSFLELL*YENKSPE
						DPQVLDNFLQVLIKVRDRHND
			!			VVPTMAQGVIEYKEKFGFDPFI
	1	1	:			STNIQYFLDRFYTNRISFRMLIN
						QHTLLFGGDTNPVHPKHIGSIDP
						TCNVADVVKDAYETAKMLCE
						QYYLVAPELEVEEFNAKAPDKP
	1					IQVVYVPSHLFHMLFELFKNSM
	1					RATVELYEDRKEGYPAVKTLV
						TLGKEDLSIKISDLGGGVPLRKI
						DRLFNYMYSTAPRPSLEPTRAA
						PLAGFGYGLPISRLYARYFQGD
	}					LKLYSMEGVGTDAVIYLKALSS
		1				ESFERLPVFNKSAWRHYKTTPE
						ADDWSNPSSEPRDASKYKAKQ
16613	46981	A	16713	175	465	ADD WOM GOLI NDAOK I KAKQ
16614	46982	A	16714	240	381	FFLSFPPLCQSS*NQNLLPW/ICA
10014	10,02	[``	1.0717	 		KVCSFTPEVSKTTNPLEGRNSG
				1		HI
16615	46983	A	16715	268	1524	
16616	46984	A	16716	1	378	
10010	1.0704	1' 1	10710	<u></u>	12,3	<u> </u>

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO in USSN	Nucleotide location of first		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
16617	46985	A	16717	708	987	IQLPGHRSQGQ\AGASHTDARQ
						QDPGNSHSDFVRHSHSLPCLER
		ł				QDRRSAGHNGSATVPGTEACG
						RAYKAPAGHYHSGAGQSLPAA
						SHRSFSSP
16618	46986	Α	16718	1	2052	
16619	46987	В	16719	l	606	
16620	46988	Α	16720	1	541	QHQRKQKRQLRAQQNLSWEEI
						AKEYQNEEDSLGGSRVVVCDI
						NKEMLKVGKQKALAQGYRAG
ŀ						LAWVLGDAEELPFDDDKFDIYT
						IAFGIRNVTHIDQALQEAHRVL
						KPGGRFLCLEFSQVNNPLISRLY
						DLYSF\RVIPVLGEVIAGDWKSY
						QYLVE\SIRRFPSQE\KFKD\MIE
•						DAGF\HKVTYESLTSG\IVAIHSG
						FKL*FLCLEFSQVNNPLISRLYD
						LYSFQGHPCPGRGHRWRLEVLS VPCREVSEGFRLRKEFKDHDRR
						CRLSTR
16621	46989	A	16721	526	933	CKLSTR
16622	46990	A	16721	191	468	
16623	46991	A	16723	2	677	GKVHVLERRGFHRTEMHLQKI
10023	10001	``	10723	1	0,7	LLLPVLSSSSSSDP/STLSVSALV
						RMFSSADE*SVSEASSSEPSSSFS
						RRSSCSFGSEPSFASFSFSSFPDL
						ASLSSVFLLFRAFS\SSHVM*WL
						PSSS*VGS/STSARAQLPSYVPSF
						SVKTKSGLKSALRFFPKAFCRL
						PSTMGPSSSSSPESDSALHRHLV
						ILDVWLQIKPRVRKPCCRRSHF
						RKRAQPAVRPLTFRSLLVVLR
16624	46992	Α	16724	31	187	-
16625	46993	Α	16725	215	947	
16626	46994	Α	16726	1	378	
16627	46995	Α	16727	2	7567	GGGGARRQRRGDAGGGAETAP
						SEIILSPAAEERSGRSETLRRGT
		1				AAGRMATVVVEATE\PEPSGSI
						ANPAASTSPSLSHRFLDSKFYLL
						V\VVGEIVTEEHLRRAIGSIELGI
						RSWDTNLIECNL\DQELKLFVSR
						HSARFSPEVPG\QKIL\HHRSERF
						*ETV\VLIQ\PFLMEGSSGTEVRL
					1	MITDAARHKLLVLTGQCFENT
1					1	GELILQSGSFSFQNFIEIFTDQEI
16600	46006	1.	1.6722	1101	265	GELLSTTHPANKASLTLFCPEE
16628	46996	Α	16728	101	365	VITGQPNPTNPNLQATPDIQGTL
			1			ARLGPA/LHPRGIIPFIGHQSA/R
					1	PHAPAPANPNKRTLFRPPPLSPP
	<u> </u>		L	<u> </u>		SPEPPGSGQGAGLGARPALP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
-						
16629	46997	Α	16729	1170	2847	VASC*ENACKQNQRQKPCDYL
						NRCRKGLWQNSTTLHAKNSQ*
						TVLYKTIKFQLEKTTLKFIWNQ
						KRAHIAKSILSQKNKAVGITLPD
						FKLYYKATVTKTAWYWYQNR
						DIDQWNRTEPSEIMPHIYNHLIF
		l				DKPDKNKKWGKDSLFNKWCW
		l				QNWLAICRKLKLDPFLTPYTKI
						NSRWIKDLHVRPKTIKTLEENL
						GNTIQDIGTGKDFMSKTPKAM
						ATKDKIDKWDLIKLKSFCTAKE
						TTIRVNRQPTKWEKIFTT
16630	46998	A	16730	79	536	
16631	46999	A	16731	232	771	
16632	47000	Α	16732	3	339	
16633	47001	A	16733 16734	279	416 1760	
16634	47002 47003	A A	16735	309	1718	
16636	47003	A	16736	10	417	IAIMNDTVAIRTRKFMTNRLLE
10030	47004	<u> </u> ^	10730	110	417	RKQMVIDVLHPGKATVPKTEIL
						EKLAQMYKTTPDVIFVFGFRTH
						CGGGKTTGFGMIYDSLDYAEK
						NEPKHRLARHGLYEKKKTS*K
		1				QRQERTNRMKKVRGTAKANV
						VAG*KPNE
16637	47005	A	16737	96	609	EKFMT\NRLTSGETKWVI*CPFT
10057	17003	`	10,3,			PGRRTVP*AQKIR\EKLAQNVQ
						RPHPDVIFVFGFRTHFGGGKTT
						GFGMIYDSL\DYAKEKMNPKH
l						RLA\RHGPGMEEGKRPSKKGN
i						GKER\KNQN*RKVQGGLAKGP
						LLGAGOKEEMKCLGSELEIGSO
		1				PKELKVLPMMLAVATVDFSQE
16638	47006	В	16738	1	1041	
16639	47007	Α	16739	3	447	VSVPFIFIWLTCKSFLKFDLKSN
						TNVLQSL/GKFQSEVSKRQKLA
						TVFSSFTSTGLCQGSVAFISRCP
						PLASGSDSVWRGGSVFFSFRP*
						RLMISLLGSCVSHQFPEKLEF**
		1				KKMPYKIPTAHHQQQDDQQRG
						HEHKQSQPDAVPHLG
16640	47008	Α	16740	1	2385	_
16641	47009	Α	16741	1	639	
16642	47010	Α	16742	1	993	
16643	47011	Α	16743	127	426	CTDKITLVFGV*MLTPKPDK/CD
						DSAV*ELLWLTNQLSACVVHP
						GLYHLHVRSIWIPYRCLFSTMC
						GVCQTCTVAMQLDCLADSFNA
						VFKLTEAAGHMAR

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
16644	47012	A	16744	3	374	SLRLQFYYKISFQTQDQANKLS DSDASHLHPLAPCSEPTRTNVH TKLPRERPQSQIASPYPLPPCPE RLKKQTNKKKNW*EPSPHILPP RNAAAHTGQSRASKSLPAPPPT VSPRPRPGAPGT
16645	47013	Α	16745	16	253	
16646	47014	Α	16746	163	588	TFHAVLHIENAGYFNSIPFFRIL NHNPLTTVEDPYLFKLPALKYL *VL*YSHESRDDLYSF*IFHQRL SILHLG*NVII\KF*LSY*KKVIGE EKN*EDV*WSRQPAGEREQC*R THIDLE/PCRHMEEYYLTKKAK GKRCSLF
16647	47015	A	16747	2	363	GTSQTLLRSLASACIFQPKVRLL SREEIPLNLNY*SVSPNSNPNHL YSRHTILTAMLTLQLGTCVLLPI GLLVPYPNCPSPTGLQVTPFLPS PCGSGWPASFTYSSNRFTTNLL FCVFIL
16648	47016	Α	16748	1028	1621	
16649	47017	A	16750	1	112	LGNTWG*QPCKRLKIWLSLEFT KINVIRHMWKKFKRL
16650	47018	A	16751	91	597	LKNRRRSRPSIRQSIGSTSVSRW LTSLFTYLDHTADVQ*V*REFIP LKPRQ*ED*MFQSWLHAWGDT LEEAFEQCAMAMFGYMTDTGT VEPLQTVEVETQGDDLQSLLFH FLDEWLYKFSADEFFIP\GWGEE FSLSKHPQGTEVKAITYSAMQV YNEENPEVFVIIDI
16651	47019	A	16752	226	668	IYFFRLHAWGDTLEEAFEQCA MA\MFGYMTDTGTVEPLQTSRS \KTQGDDLQSLLFHFLDEWLYK FSADEFFIPREVKVLSIDQRNFK LRFN\GWGEEFSLSKHPQGTEV KAITYSAMQVYNEENPGSFCD HWTFKTTQKIKRLPTGKK
16652	47020	Α	16753	3	942	
16653	47021	A	16754	3	419	SLYHNSSQKRHWTFSSEEQLAR LRADANRKFRCKAVANGKVLP NDPVFLEPHEEMTLCKYYEKRL LEFCSVFKPAMPRSVV\LTCAFL ACKVDEFNVSNPQFVGNLRESP LGQEKALEQILEYELLLIQQLNF HLIVHN
16654	47022	Α	16755	312	537	
16655	47023	В	16756	209	1381	
16656	47024	A	16757	1	927	
16657	47025	Α	16758	1	3987	
16658	47026	Α	16759	2	175	

SEQ ID	-		SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
16659	47027	Α	16760	2395	2700	
16660	47028	A	16761	2	1118	
16661	47029	A	16762	26	1205	
16662	47030	A	16763	1	1594	
16663	47031	A	16764	1	595	
16664	47032	Α	16765	1 .	1029	
16665	47033	A	16766	1172	1364	
16666	47034	A	16767	253	389	
16667	47035	Α	16768	489	855	
16668	47036	Α	16769	2	401	
16669	47037	A	16770	415	694	EAKLHLNICDMSWLCCPTKGK RKQALEVLNILGPFLAPFWIAK SYFFTSPKTQVSSL*FLRYKRIK LQCRPKTCTNPSRTRSCALPGR FIPPF
16670	47038	Α	16771	1	765	
16671	47039	В	16772	1	1095	
16672	47040	A	16773	82	311	KANSDVPFGFGVNLEHLMEAC NS/GTNLLNSKRLNNDCICLASH TDVSIIARKCCLVVSCLLLLRGL SILRSKESHC
16673	47041	Α	16774	2	683	
16674	47042	Α	16775	1	2703	
16675	47043	Α	16776	1	555	DHNSSPAREQ/SSMENKFHELT EVGFRKITSLEKNINDLMELKN TARELREAYTSIKSRIDQAKESI SEIEDQLNEIKCEDKVREKRMK SNKQNFQEIWDYVKRPNLRLT GVPESDRENGTKLENTLQEIIQE NFPNLARQANIQIQEIQRTAQR QSSSRATPIHIIVRFAKFEIKEKM LRAAREK
16676	47044	Α	16777	1	839	· · · · ·
16677	47045	A	16778	1	504	
16678	47046	A	16779	1	2367	
16679	47047	Α	16780	1	1566	
16680	47048	A	16781	146	464	VTGWTNYEGNGRWRERSRQLC GCCSRKSSSSWGISLMKGSGAP LRPGRMMWSGFRKCSDTQVM YS*R*LLETMTLASIMR*THTK* NALRKCSALKDCFLGKALTL
16681	47049	Α	16782	1	1782	
16682	47050	В	16783	221	1075	
16683	47051	Α	16784	1	627	
16684	47052	Α	16785	1	1539	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
	of peptide	1	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
16685	47053	A	16786	1	1318	LSPVLSLSPDSMSFTTRSTFSTN
10002	17023	'	10700		1310	YRSLGSVQAPSYGARPVSSAAS
,		l				VYAGAGGLATGIAGGLAGMG
						GIQNEKETMQSLNDRLASYLDR
						VRSLETENRRLESKIREHLEKK
						GPQVRDWSHYFKIIEDLRAQIF
						ANTVDNARIVLQIDNA\RLAAD
						DF\RVKYETELAMRQS\VENDIH
						GLRKV\IDDTNITRLQLETEIEAL
			<u> </u> 			KEEL\LFMKKNHEE\EVKGLQA
				Ì		QIASS\GLTVEV\DAPKF\QDLRQ
						DSWADIR\AQY\DELARKEPRG
						A*TKYWFSADLRESTTV\VTTQ
						F\AEV\GAA*DDASQSWRTYSPS
		!				SLGDPTWGFPWRKS*RAS\LEN
						SLREG*RPAYALT/QMEQASTGI
						LACTFESELGTDPGAEGQRQAQ
		:				EY\EALLNIKVKLEAEI\ATYRRL
						\LEDGEDFNLGDALDSSNSMQT
						IQKTTTRRIVDGKVVSETNDTK
16686	47054	A	16787	1	253	RVRTLKTSTSTCLYWFENKDS/
10080	47034	\^	10787			DHPERSSP*YMSPSHRYGSLVK
						ASITKPAFLKTSGRLFTCSSQFC
		1				PLISKISDGFSHLPPFST
16687	47055	A	16788	1	276	AAAAGPPGPLVMFTYRERVSG
10007	1,033	ľ.,	10700		[· · ·	LRSVHILRASHTDARLRQEEPT
						ERLQFHSSGLLIRFSRETEPIGV
						YIKED*EEWAYKMMVFAKSHD
	i					LPFAN
16688	47056	A	16789	1	203	LLHTSTKNELWQ*GKRDPLIVF
						YTQYLF*EKTTRK*NQRQAGSL
						ALGPKPGLRLPGLNPVVENQFI
16689	47057	A	16790	8	361	RGLDGQGGRDGSSSGGGGGGR
		1				RRRCLLLLAPAPAVIRASGRGP
						R*DCWRAGCA*AWRPC/RLAR
						RMWTLRSPLTRSLYVNMTSGP
						GGPAAAAGGRKENHQVARKR
						LNVCVFTRSVSETA
16690	47058	A	16791	674	1076	VQLFRSAGSRDHHRFLGRGRN
						KQTKTLYPTPIPCSLKYQRKQN
						GSLLWTSRMPSSVFPCTLTPSFS
						LPLRISQTTHPNLRGPSCPKGLG
						IALISLVRHWLKI*ATSQVQAL
						WSFSMWMIYFWLPVQKPHASR
16691	47059	В	16792	296	1470	
16692	47060	A	16793	1	334	
16693	47061	Α	16794	718	2541	
16694	47062	A	16795	2	406	
16695	47063	Α	16796	74	464	
		_			610	
16696	47064	Α	16797 16798	159	610 273	

SEQ ID NO:	of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
16698	47066	Α	16799	1	543	
16699	47067	В	16800	1	873	
16700	47068	Α	16801	1	558	
16701	47069	Α	16802	27	136	
16702	47070	Α	16803	89	363	
16703	47071	Α	16804	579	994	
16704	47072	A	16805	129	362	PKIIQPRRATTRFTSIHGWRICPS IPCFPEATINRRRCVKVDSVTCH LLTQR*YKSSKVQERC*SCRIQG QKISDCY
16705	47073	Α	16806	878	1811	
16706	47074	Α	16807	l I	843	
16707	47075	С	16808	19	231	
16708	47076	A	16809	1	1650	
16709	47077	Α	16810	3	404	
16710	47078	Α	16811	279	428	
16711	47079	A	16812	3	507	
16712	47080	Α	16813	1	1128	
16713	47081	В	16814	1	567	
16714	47082	A	16815	3	1206	GSEGQGAGPGGAAATAAAME DEMPKTLYVGNLSRDVTEALIL QLFSQIGPCKNCKMIMDTAGN DPYCFVEFHEHRHAAAALAAM NGRKIMGKEVKVNWATTPSSQ KKDTSNHFHVFVGDLSPQITTE DIKAAFAPFGRISDARVVKDMA TGKSKGYGFVSFFNKWDAENA IQQMGGQWLGGRQIRTNWATR KPPAPKSTYESNTKQLSYDEVV NQSSPSNCTVYCGGVTSGLTEQ LMRQTFSPFG\QIM\EIRVFPDKG Y\SFVRFNS/HMESGRHMAIVSV NGTTIEGHVVKCYWGKETL\D MINPV\QQQNQIGYPQPYGQWG Q\WYGNAQQIGQYMPNGWQV P\AYGMYGQAWE/SSQGFNSDT VFLAPWDGDPNFGSCNRLQGQ NG\SLLPN\RPSGYRVAGYETQ
16715	47083	Α	16816	1425	1564	
16716	47084	С	16817	151	339	
16717	47085	Α	16818	67	373	FCDCHHFILMFKSPHIWPVGIFS SWLLCFFWACLHHSLSIALLSC TKRYSGLILYFLCSSFEITVSSKS SVSF*RRMVFRNQVLGSRCACC C*GVAAPRPFP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
16718	47086	A	16819	1	799	MGPHWLWLHLALVLLPPSALL
						LPSCEKGVLRVRFYHEKWIYV
						HKESTKERHGYCTLGEAFNRL
		ļ				DFSSAIQDIRRFNYVVKPLLPRY
		1				KFQRLNAITKDRQRSSSGVQLP
		ŀ				HAGQPQGADPTQDERRQVAGD
						TPGDPRALAGGGVVDRTYPGS
						YEASGRPGLQLALAIEDLLRDS
						AIGCTFCDRCRPVGAQRLLAHS
		l				LALAAPGAEKREVVKLLLVAY
		Ì				VRRRPGKNLTTQNCQ\PAAGYI
1		l				PPGHQ*AASAHARSWKŅQQPP
						PRVLIGSA
16719	47087	В	16820	28	438	
16720	47088	Α	16821	366	793	A WVEQSKVLIKEGGIQLLLTIV
						DTPGFGDAVDNSNCWQPGINYI
						DSKFEDYLNAESQVNRCQMPG
				1		NRVHCCLYFIAPSGHGPLHN*R
						LPPSGRIG*YMFVTTWHCLLLR
						LKPLDIEFTKHLHEKVNIIPLIAK
		<u> </u>				ADTLMPEEC
16721	47089	C	16822	228	374	A WEY SWOWN DD GI DDGID ID WD
16722	47090	Α	16823	291	604	A WELEKSHLRRGLDPSHNINIR
						VEPKF*SKYGGVQLLLTIVDTP
						RFGDAVDNSNCWQPVINYIDIK
·	1	1				FEDYLNAESRVNRCQMPGNRV
16500	15001	 	1.600.4	140	400	Q**LYFIAPSGHGPLHN
16723 16724	47091	A	16824	148	498 595	DRPRITGGGKSGTEYPENLPTL
16/24	47092	A	10823	1	393	KATIENKNSVLNTATKMKDVQ
						TSTPAEQDLEMA\SEGEQKRLE
						EYENNOPOVKN\QIHSRDDLDD
						IIQSSQTVSEDGD\SLCCNCKNVI
		1				LLIDQHEMKCKDCVHLLKIKNT
						FCLWKRLIKLKDNHCEQLRVKI
						RKLKNKASVLQKRISEKEEIKS
						OLKHEILELEKELCSLRFAIQQE
16725	47093	A	16826	1	828	Z=====================================
16726	47094	A	16827	1	540	
16727	47095	Α	16828	295	460	
16728	47096	Α	16829	2	1355	
16729	47097	Α	16830	650	1184	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
16730	47098	A	16831	725	2368	AVDIIKEIKHPQKHKKLPAFQLR PFPLREETIPCNGVSFPRCEPPCL AEVYMCLNEMLKDEKKGCMN KSGAGENPQERTAQKRKFPSPP HSSNGHSPQDTSTSPIKKKKKP GLLNSNNKEQVKQPLECIETQS KAMTMLTIEQLSYLLKFAIQKM KQPGVRRKLKPVPLEQHPDYA EYIFHPMDLCTLEKVGLRQMY GCTEAFLADAKWILHNCIIYNG GEWLRTQIAKVVIKICEHEVCV SQVCPECYLAACQKRDNWFCE PCVSWIYD\SACQPV*WKLVIQ MFLPM/WKADIL/RGEEG*YTSE LIGWL*SLSCDCSAGTSISHPSD SD*RWLTEHEGRRVSWP\VNNC YLMSKEIPFSVKKTKSIFNSAM QEMEVYVENIRRKFGVFNYSPF RTPYTPNSQYQMLLDPTNPSAG TAKIDKQEKVKLNFDMTASPKI LMSKPVLSGGTGRRISLSDMPR SPMSTNSSVHTGSDVEQDAEK KATSSHFSASEESMDFLDKSTG QLRAGERGGGPLDTGVSHGCT LGGPKQVSIKQQASVLQDTMN
16731	47099	A	16832	2	164	PSTLRAEAWPP*NGSPVPLASS QASSSRSGPTSCIPRYPVASPSE RLPLPTAAA
16732	47100	Α	16833	51	498	
16733	47101	Α	16834	1	531	
16734	47102	С	16835	257	358	
16735	47103	A	16836	63	193	SGRLAPHTSRRTSANCSDDAKS SDSCSPSRKT*WSGRNTNRIH
16736	47104	Α	16837	1	834	
16737	47105	A	16838	258	545	CQNPQILASPSTKDLMGLGRNT NKDTLKACLARVVKRRSECSQ KPAS*MCTERNPTSMREVHDLE RKYAVLYQPLFD*AVLKLLMQ FMNLRKEGM
16738	47106	Α	16839	66	476	
16739	47107	Α	16840	1	1362	
16740	47108	A	16841	1	747	
16741	47109	Α	16842	522	687	
16742	47110	Α	16843	112	861	
16743	47111	A	16844	87	472	TVTFSQCRFGKMLPLEKAFASP QELPSPAGSAHAGVSSRSSSRK NPRPSLRGPLLTLEFSRLRFREF VYP\EAAGPHHTLARLDELCRQ WLMPEARSKEQMLELLVLEQF L/SILPDKVRPWVVAQYPES
16744	47112	Α	16845	1	393	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
16745	47113	Α	16846	621	1099	
16746	47114	Α	16847	1	453	
16747	47115	A	16848	209	1175	
16748	47116	Α	16849	1	546	
16749	47117	Α	16850	187	421	LGESSQ*GSP\PPPTGGCGPATG
						Q*PLGQSFQRKELAAIFAVSQY
						SLVVPPGMGETEATGIWSRPPA
						NGNSPTVEWHIK
16750	47118	Α	16851	3	1654	
16751	47119	A	16852	166	465	
16752	47120	Α	16853	1	558	
16753	47121	Α	16854	51	380	RNNLLLLTHHFVLIIQFLIPRKET
						KYYLVITPYSLSINVLSNFFYYS
						PESLNDTNCYHHEEADTPWR*F
						LAIFEGLFNIAHQMQNMPFFLL
						WKQVPETLVIQQQLPRIN
16754	47122	Α	16855	3	550	
16755	47123	A	16856	1	462	
16756	47124	В	16857	1	2811	
16757	47125	A	16858	3	514	ASGTAPEIQARPTRPRRKRRK
i	! !					LLPL/TKAEAKAKALKAKKAVL
						KGVHSHKKKKIRTSPTFRRPKT
		ŀ				LRLRRQPKYPRKSAPRRNKLDH
						YAIIKFPLTTESAMKKIEDNNTL
						VFIVDVKANKHQIKQAVKKLY
						DIDVAKVNTLIRPDGEKKAYVR
						LAPDYDALDVANKIGII
16758	47126	A	16859	73	396	FFQFSPKKNCSVWWNC\AEVLL
						KYKTGETNDFELLKNQLLDPDI
		i				KRLPWLNRSQTVVEEYLAFLG
						NLVSAQTVFLRPCLSMIASHFV
						PPRVIIKEGDVDVSDSDDEDD
16759	47127	A	16860	132	357	YVSKFAGRLSSSSSESETSTSPS
						LMITRGGTKWGSNHAETRSEE
						NSLC*YKITKKSQILFHYCFDFY
		ŀ				STKAISI
16760	47128	Α	16861	380	597	
16761	47129	A	16862	179	280	QRDIKSCVQILQWEGSTRPFTN E**YMANVRNS
16762	47130	A	16863	2	175	
16763	47131	A	16864	178	374	
16764	47132	A	16865	386	2120	
16765	47133	A	16866	887	1104	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence		in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
16766	47134	A	16867	1	708	MEVAALEHLDHRLREKVKQQN ALRHQVVLRQRRLEELQLQHS
ł						LRLLEMAEAQNRHTEVAKTMR
		ļ				NLENRLEKAQMKAQEAEHITS
						VYLQLKAYLMDESLNLENRLD
						SMEAEVVRTKHELEALHVVNQ
						EALNARDIAKNQLQYLEETLVR
						ERKKRERYISECKKRAEEKKLE
						NE/LHGAQADPGSDLNPRADPP
						RAPAATVRRHHPGQPACQGGG
						AAAALEHVPDGGDLWQGQGR
	,	<u> </u>				HWH
16767	47135	A	16868	11	1327	
16768	47136	Α	16869	1	1776	
16769	47137	A	16870 16871	1	397 1974	
16770 16771	47138 47139	A	16872	1	336	
16772	47140	A	16873	3	661	GRGGHNFAPNLTARSAVTSGL
10772	77140		10075	ľ		GGPPAAVMVGSLNCIVAVSQN
						MGIGKNGDLPWPPLRNEFRYF
		1				QRMTTTSSVEGKQNLVIMG\KK
						TW\FSIPE/RRNRPFKG*EFNLVL
						SRELQGNLPQGAHFLSRSLDDA
						LKLTEQPELANKVGMVW\IVG
		1				GSSI\YKESMNHSSSLKLFVTRII
						QDFESDTFFPEIDLEKYKFLPEY
		<u> </u>				PGVFFDVQEEKGIKYKFEVHEK
16773	47141	Α	16874	38	755	ALKTCSVNQPDAKDNIAGSSGH
						LRTLSPVVSRSCLKEEERRRRR
						RMKRRQRSRRRRSRRRRMRRR
						RRSRSRRRRMKRRKRRRMKR
						RSSFILMYFFMLLLSGIPLLYME VIMGQWLHVDNIRVWKQLVP
						WLCSMSY/SSQSVCASVSLYNS
						TIISWNFFYSFAHPLPWDHCPLV
						KNISVTDWAHQYFLYHTTLHA
						SDHSEEAAEALVPNRSLGCLLG
						RDHRDLDFKPEIRRGAHTGGAR
16774	47142	Α	16875	1	1830	
16775	47143	Α	16876	2	1142	
16776	47144	Α	16877	1	435	
16777	47145	Α	16878	1	963	
16778	47146	В	16879	247	1092	
16779	47147	Α	16880	1	1653	
16780	47148	A	16881	1	2268	
16781	47149	A	16882	301	555	
16782	47150	A	16883	1	630	
16783	47151	A	16884	731	834 2277	
16784 16785	47152 47153	A	16885 16886	45	2446	
16786	47154	B	16887	625	696	
10/00	14/134	טו	1,0001	1023	1020	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
16787	47155	A	16888	193	315	WPGGGYSLLCFWCPLSFAGAA
10,0,	' ' ' ' ' ' ' ' ' ' ' ' ' ' ' ' ' ' '		10000			G/W*PPPGHH*PYSPKQEHYRT
						NSDTLIFLSSRTLVLQGS*GIKV
						AGSEGPKYNE*PRNPVLPSLCL
		İ				HLFGC
16788	47156	A	16889	1	588	
16789	47157	Α	16890	2	229	
16790	47158	A	16891	1	502	
16791	47159	Α	16892	516	576	
16792	47160	Α	16893	144	435	
16793	47161	С	16894	1	996	
16794	47162	Α	16895	424	632	
16795	47163	В	16896	446	1944	
16796	47164	С	16897	207	470	
16797	47165	Α	16898	1	396	
16798	47166	Α	16902	1	585	
16799	47167	Α	16903	141	906	
16800	47168	В	16904	99	276	
16801	47169	Α	16905	1	756	
16802	47170	Α	16906	1	383	
16803	47171	Α	16907	1	582	
16804	47172	Α	16908	3	615	PECIIGIDILSSGQNPHIGSLTGR
1						VRAIMVGKAKRKPLELPLPRKI
						LNQKQYRIAGGIEEISATIKDLK
						DAGVVIPTTSLFDSPIWPVQKT
						DGSWRMTVDYRQINQVATPIA
						AAIPDVASLLKQINTSPDTW/PI
ŀ						RPPISNGD*GVSGR*ACCLEPLA
						GPHR*ITSEASRILEQGPAIFCR*
						LLSF*ETALGLLLGFGGN*TFDY
16805	47173	Α	16909	3	415	PVK/VGA*GGQVINGVLAQV*L
						TVGPVGPRTHPVVIFPVPECRIG
						RDILSSWQNPHTGSLTGRVRAF
						MVGKAKWKP*ELPLPRKTVNQ
						KQYRIPGGIAEIS/A/TIKNLRRG
		1				VVIPTTSRFNSPIWPVQKTDGS
						W*TAADY
16806	47174	Α	16910	232	633	
16807	47175	Α	16911	1	1677	
16808	47176	Α_	16912	328	381	LLKLQH*MQSPYLVGPSQ
16809	47177	A	16913	3	811	
16810	47178	A	16914	-	1125	
16811	47179	A	16915	1112	1191	
16812	47180	A	16916	113	485	
16813	47181	A	16917	1	357	
16814	47182	A	16918	12	398	TENA A COMPONIO DI IN ESTAT PERM
16815	47183	A	16919	498	595	TEMASTTTCT/RFHDEYQLFEEL *KGAFSVNGT
16816	47184	Α	16920	5	367	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
16817	47185	A	16921	190	2005	VLYPLHLGTGTGKGAGGASSG EEGSEARRRRIGGLWGRRPHA TLSGHLPSRPAPAHTPSRASYFR TRKVRAALGDSAAGPVPGLAA RLLASGPSRSQPGHSPGPEMAS TTTCTRFTDEYQLFEE\LGRGAF SVVRRCMKIPTGQGYAAKIINT KKLSARDHQKLEREARICRLLK HPNIVRLHDSISEEGFHYLVFDL VTGGELFEDIVAREYYSEADAS HCIQQILESVNHCHLNGIVHRD LKPENLLLASKSKGAAVKLADF GLAIEVQGDQQAWFGFAGTPG YLSPEVLRKDPYGKPVDMWAC GVILYILLVGYPPFWDEDQHRL YQQIKAGAYDFPSPEWDTVTPE AKDLINKMLTINPAKRITASEAL KHPWICQRSTVASMMHRQETV DCLKKFNARRKLKGAILTTML ATRNFSAAKSLLKKPDGVKEST ESSNTTIEDEDVKARKQEIIKVT EQLIEAINNGDFEAYTKICDPGL TAFEPEALGNLVEGMDFHRFYF ENALSKSNKPIHTIILNPHVHLV
						GDDAACIAYIRLTQYMDGSGM PKTMQSEETRVWHRRDGKWQ NVHFHRSGSPTVPIKPPCIPNGK ENFSGGTSLWQNI
16818	47186	Α	16922	1	288	
16819	47187	Α	16923	318	458	
16820	47188	Α	16924	1	578	
16821	47189	Α	16925	803	1068	
16822	47190	Α	16926	1	660	
16823	47191	A	16927	2	325	TSSENNNGPKR*IHSTESLYLLQ GSVVADHDYIGLPEIPIGAYQA NILVEDATIGIVDNELLTSSKDR ELLEYRNTKISPLIDDHSSLEKQ TFSLLDSSNQVLEYLS
16824	47192	Α	16928	156	573	
16825	47193	A	16929	405	599	GLEPPQKPGLVTSILQANPVPLP CGNSQSR*TSTRCWYEHLQCTQ SHLQMQQYHEAYQSACALA
16826	47194	В	16930	1	1676	
16827	47195	Α	16931	1	1125	
16828	47196	Α	16932	27	818	
16829	47197	Α	16933	1	4986	
16830	47198	Α	16934	1	4401	
16831	47199	A	16935	153	370	SVLETCMSTAFPVSCHSSFKTA VLRISLS*ISAQGGPIDLFSQCHP SSSTGVSS\NHRGSWDTPGQSL VVICL

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
16832	47200	В	16936	47	501	
16833	47201	Α	16937	49	1714	
16834	47202	A	16938	1	999	
16835	47204	A	16939	31	2029	FAARAGPGGSRRARRGNKWAR MPYEIKKVFASLPQVERGVSKII GGDPKGNNFLYTNGKCVILRNI DNPALADIYTEHAHQVVVAKY APSGFYIASGDVSGKLRIWDTT QKEHLLKYEYQPFAGKIKDIAW TEDSKRIAVVGEGREKFGAVFL WDSGSSVGEITGHNKVINSVDI KQSRPYRLATGSDDNCAAFFEG PPFKFKFTIGDHSRFVNCVRFSP DGNRFATASADGQIYIYDGKTG EKVCALGGSKAHDGGIYAISWS PDSTHLLSASGDKTSKIWDVSV NSVVSTFPMGSTVLDQQLGCL WQKDHLLSVSLSGYINYLDRN NPSKPLHVIKGHSKSIQCLTVH KNGGKSYIYSGSHDGHINYWD SETGENDSFAGKGHTNQVSRM TVDESGQLISCSMDDTVRYTSL MLRDYSGQGVVK\LD\VQPKC\ VAVGPGGYAVVVCIGQIVLLK DQRKCFSI\DNPGYEPEVVAVH PGG\DTVAIGGVDGNVALYSIL GTTL\KDEGKLLEAKGPVTDVA YSH\DGAFLAVCDASKVVTVFS VADGYSENNVFYGHHAKIVCL AWSPDNEHFASGGMDMMVYV WTLSDPETRVKIQDAHRLHHVS SLAWLDEHTLVTTSHDASVKE
16837	47205	A	16941	1	53	
16838	47206	A	16942	1	389	GMPTSTIASCSQDGRVFIWTCD DASSNTWSPKLLHKFNDVVWH VSWSITANILAVSGGDNKVTEG QQNEQ*QDRWGLAPHPPAPGL PLPGPTNQTTGKSPQLQQDYFP RRSYRCSHRLIICLNVIGNAL
16839	47207	A	16943	1	438	
16840	47208	В	16944	1	1267	
16841	47209	A	16945	635	808	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hođ	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
16842	47210	Α	16946	37	1319	QGEGHGRGLHGFLGRPESQIPS
						WSRSAPLHRLEIWGALAPSLAD
						HSEGSPEVSVINTVDTSHEYMI
		l				Y\DAQMDYYGHPPGQPASSDRS
		1				VKIF*CAQWEGRSLFADLQGS
						WRGPVWQVGWGFTPLLRATS
]				WAFVASLLTGKFIYSWERGKT
		1				GTLGRRATSNAGTRLLKLNFGC
						CWAPHGLTVLHSWACWVVSD
			:			RGPSSL\LDFTPGEGQWEVKKIN
		l				NAHTIGCNAV\TWAPACCTWK
		l				PHRRPHRGQKPNYIKRFASGGC
ľ		1				\DNLI\KLWKEEEDGQWKEEQK
		ŀ				LE\GHSDWV\RDVAWAPSIGPA
						HQHHRPAAPRMGRVSIWTCDD
						ASSNTWSP*IV\DKFNDVVWHV
						S\WSITANILAVL\CGDNK\VPL
						WKESS*WAKWVVPSSDVKQRG
						PGLPYQPSSDQEGPARNEQVNK
						DQGGALAPPPAQLPGPAPSPGP
16843	47211	Α	16947	1	1021	MSGEFIDHSHIARIIYPGIFGTDA
			1			GSCIDQRGITGKLQQPPEDRRL
						GEGKLTNTKDIHTKTPSVRHHH
						QRPKVDKTTKIGKKQSRKTGNS
						KNQSAPPPPKECSSSPAMEQSW
						TENDFDELREEGFRRSNYSELK
						EEFRTHGKEVKNLEKRLDEWL
						TRITNAEKSLKDLMELKTKARG
						LQRVSVMEDEMNEMKQEEKFR
		l				EKRIKRNKQSLQEIWDYVKRPN
						LRLIGVPESDGENGTKLENTLQ
						DIIQESFPNLARQANIQIQEIQR
1		ł				MPQRDSLRRATPRHIIVRFTKV
		l				EMKEKMLRAAREKA\HIALIPK
		i				LTT*LEVKLSSANVKEQKL*QT
						VFQTTVQSN
16844	47212	A	16948	1	393	
16845	47213	A	16949	1	2631	
16846	47214	A	16950	1	1155	
16847	47215	A	16951	1	1185	
16848	47216	A	16952	l .	942	
16849	47217	A	16953	1	615	

SEQ ID NO:	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO.	sequence	nou	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
16850	47218	Α	16954	474	2678	IKPQRWGKKQSRKTGNSKNQS
					,	ASPPPKERSSSPAMEQSWTEND
		Ì				FDELREDSFRRLVITNYSELKEE
					}	VRTNGKEVKNLEKKLDEWLTR
						ITNAEKSLKDLMELKTMAQELP
		1				DQCTSLSSRCDQLEERVSVMED
						EMNEMKCEEKFREKRIKRNEQ
						SLQEIWHYVKRPNLRLIGVPES
						DGENGTKLENTLQDIIQENFPN
						LARQANIQIQEIQRMPQRYSSR
						RATPRHIIVRFTKVEMKEKMLR
1					:	VAREKGQVTHKGKPI\RLTADL
		ļ				LAETLQARREWG\PIFNILKEKN
				İ		FQPRISYPAKLSFIDRSTRQKVN
						KDTQELNSALHQADLIDIYRTL
	l	ŀ				HPKSTEYTFFSAPHHTYSKIDHI
		į				VGSKALLSKCKRTEIITNCLSDH
						SAIKLELRIKKLTRNHSTTWKL
						NNLLLNDYWVHNEMKAEIKM
						FFETNENKDKTYQNLWDTFKT
						VCRGKFIALNAHKRKQERSKID
		1				TLTSQLKGLEKQEQTHSKASRR
						QEITKIRAELKERVGRWRIGEA
						ADLVGVSSQAIRDAEKAGRLPH
	ŀ					PDMEIRGRVEQRVGYTIEQINH
						MRDVFGTRLRRAEDVFPPVIGV
						AAHKGGVYKTSVSVHLAQDLA
						LKGLRVLLVEGNDPQGTASMY
						HGWVPDLHIHAEDTLLPFYLGE
						KDDVTYAIKPTCWPGLDIIPSCL
	İ					ALHRIETELMGKFDEGKLPTDP
						HLMLRLAIETVAHDYDVIVIDS
16851	47219	A	16955	1	675	
16852	47220	A	16956	1	1032	
16853	47221	Α	16957	3	978	HEAKHQMADDAGGSGGPKA
						LVGPGMG\NPVAFRRGF\GIVIR
						GPGSRPGGRGRGR\GRGARGSK
ļ						\AEDKDW\MPVTKLG/RALVKD
						H*RSKFPWKEIYLFSLPH*RNQR
						IIDF\FLGGLLSKDEGFE*LCPVQ
		1				EQDPCPASRTQASRPFVAIGGL
		-				QMAHVGLGC*VLPRKVATGHP
						VGAIILAKLS\IVPVRR\GYLGNK
						VLAKP\HTVP\CKV\TGRCGSVL
						VRLIPAP\RGLGMVSAPVA\KKL
						LMMAGIDDCYT\SARG\CTATL
						GQIWPRATL*LPFLRTYKLP*PP
						DLWKGRLYLPSFPIKEFT\DHLV
						KDPHPESSVQQDSELQLVATT
16854	47222	В	16958	124	274	
16854	47222	В	16958	124	274	KDI III ESSYQQDSELQLYAI I

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
16855	47223	A	16959	202	2264	KWPKMNPQQQRMAAIGTDKE LSDLLDFSAMFSPPVNSGKTRP TTLGSSQFSGSGIDERG\GTTSW GTSGQPSP\SYDSSRGFTDSPHY SDHLNDSRLGAHEGLSPTPFMN SNLMGKTSERGSFSLYSRDTGL PGCQSSLLRQDLGLGSPAQLSS SGKPGTAYYSFSATSSRRPLH DSAALDPLQAKKVRKVPPGLPS SVYAPSPNSDDFNRESPSYPSPK PPTSMFASTFFMQDGTHNSSDL WSSSNGMSQPGFGGILGTSTSH
						MSQSSSYGNLHSHDRLSYPPHS VSPTDINTSLPPMSSFHRGSTSS SPYVAASHTPPINGSDSILGTRG NAAGSSQTGDALGKALASIYSP DHTSSSFPSNPSTPVGSPSPLTG TSQWPRPGGQAPSSPSYENSLH SLQSRMEDRLDRLDDAIHVLRN HAVGPSTSLPAGHSDIHSLLGPS HNAPIGSLNSNYGGSSLVASSR SASMVGTHREDSVSLNGNHSV LSSTVTTSSTDLNHKTQENYRG GLQSQSGTVVTTEIKTENKEKD ENLHEPPSSDDMKSDDESSQKD IKVSSRGRTSSTNEDEDLNPEQ KIEREKERRMANNARERLRVR DINEAFKELGRMCQLHLKSEKP QTKLLILHQAVAVILSLEQQVR
						ERNLNPKAACLKRREEEKVSA VSAEPPTTLPGTHPGLSETTNP
16856	47224	Α	16960	1190	1327	
16857	47225	A	16961	386	554	ATISFRYSQCK*AEQGCDSELG LFTKIRKSGYWGV*ATISFRYSQ CKALEIHLTAPVVPSALRIWGF QIPGFNQLSFHPSPESKGSFLPC
16858	47226	В	16962	499	915	
16859	47227	A	16963	1	1785	
16860	47228	Α	16964	1	3036	
16861	47229	A	16965	2	710	CC*VDWWCCD*UDO*CTI CA DI O
16862	47230	A	16966	349	493	SS*VRWVSGR*HPQ*STLSAPLG ETSGCHQGGLHDLKIVVL
16863	47231	Α	16967	1	1335	
16864	47232 47233	A	16968 16969	93	426 345	APALCLQILLDCKSVRNRTSRF RPREVGIQQYKKGGGAAGKGG TGNAPGFLLFSKDS*KDPTPLPP PLLL*GQRDH*IQLHQE
16866	47234	A	16970	1	1302	
16867	47235	A	16971	1	1174	
16868	47236	Α	16972	274	1428	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
16869	47237	A	16973	411	629	HCLSHFQNTGPRGDVDFQQEL\ FPVLMADIRKEERSHLCRSSRR TWTILDKS*IQGSRLSSKEQGW GWGTSRK
16870	47238	Α	16974	618	1552	
16871	47239	Α	16975	1	558	
16872	47240	Α	16976	1	765	
16873	47241	Α	16977	2	418	
16874	47242	Α	16978	130	1965	
16875	47243	Α	16979	1	1062	
16876	47244	Α	16980	1	558	
16877	47245	Α	16981	37	676	
16878	47246	Α	16982	1	1014	
16879	47247	Α	16983	1	2964	
16880	47248	В	16984	259	1751	
16881	47249	Α	16985	1	987	
16882	47250	Α	16986	1	1047	
16883	47251	A	16987	3	1413	GLHAARGPWVVQAWISITM\ER KEVCYGQLGCFSDEKPWAGTL QRPVKLLPWSPEDIDTRFLLYT NENPNNFQLITGTEPDTIEASNF QLDRKTRFIIHGFLDKAEDSWP SDMCKKMFEVEKVNCICVDWR HGSRAMYTQAVQNIRVVGAET AFLIQALSVKPCLGPVLWVGVR EGQGDAARGTALEDVHVIAHS LGAHT/AAEAGRSWGPRGRITD DLGTA*YKVDGMG*ERRGVRF SI*P*GGGV*AGFGGERFR*RKN VGKSTAGWRM\GLDPAGPCFQ DEPEEVRLDPSDAVFVDVIHTD SSPIVPSLGFGMSQKVGHLDFFP NGGKEMPGCKKNVLSTITDIDG IWEGIGGFVSCNHLRSFEYYSSS VLNPDGFLGYPCASYDEFQESK CFPCPAEGCPKMGHYADQFKG KTSAVEQTFFLNTGESGNFTSW RYKVSVTLSGKEKVNGYIRIAL YGSNENSKQYEIFK
16884	47252	В	16988	95	315	
16885	47253	С	16989	1	780	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
16886	47254	Α	16990	2	1426	SGTLDRMLIFWTITLFLLGAAK
						GKEVCYEDLGCFSDTEPWGGT
						AIRPLKILPWSPEKIGTRFLLYT
		1				NENPNNFQILLLSDPSTIEASNF
						QMDRKTRFIIHGFIDKGDESWV
			•			TDMCKKLFEVEEVNCIFV\DWK
ļ						KGSQATYTQAANNVRVVGAQ
						VAQMLDILLTEYSYPPSKVHLI
						GHSLGAHVAGEAGSKTPGLSK/
ŀ						RLQGLDPVEASFESTPEEVRL\D
						PSDADFVDVIHTDAAPLIPFLGF
			i			GTNQQMGHLDFFPNGGESMPG
						CKKNALSQIVDLDGIWAGTRDF
						VACNHLRSYKYYLESILNPDGF
						AAYPCTSYKSFESDKCFPCPDQ
ŀ						GCPQMGHYADKFAGRTSEEQQ
						KFFLNTGEASNFARWRYGVSIT
İ						LSGRTATGQIKVALFGNKGNTH
						QYSIFRGILKPGSTHSYEFDAKL
						DVGTIEKVKFLWNNNVINPTLP
						KVGATKITVQKGEEKTVYNFCS
						EDTVREDTLLTLTPC
16887	47255	Α	16991	1	1407	
16888	47256	Α	16992	954	1663	EPSPDPAHAPSVALLHTGNQSK
ļ						VLPWPRNPCMKRW*K*KMLKI
						SWVHEWVTSNWISTAVRSLRV
		<u> </u>	_			FAPKNVFQ**ALGPSGWLVYC
16889	47257	A_	16993	3	242	
16890	47258	A	16994	1	1815	
16891	47259	Α	16995	1	375	
16892	47260	A	16996	3	2005	
16893	47261	A	16997	114	651	
16894	47262 47263	A	16998 16999	114	275 270	
16895		A	17000	140	876	PPALVVGRRQKSWRMFEQMRA
16896	47264	A	1 7000	140	870	NVGKLLKGIDR\YNPENLATLE
						1
1						R\YVETQAKENAYDLEPNLAVP ESCTKFNP\AFFQTNGSTAQDSC
ŀ						*KALTN\LPHT\DFTL\CKC\MIRP
						GHIQE\ERPIPTDFCYLGGPGWE TWPFSRAFWESPGMKTVDLLW
						KV*LGF\EDSV\RKFI\CHVVG\IT
						YQHIDR\WLLAEMLGDLFGQAS
						*RCW\MSKYGWSADESGQIFIC
						SPRREHLTPRNIVEKIDFDSVSSI
L	l	1	L	L		MASSQ

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
1.600-	1.50 65	 	1		l	L CONTROL CONT
16897	47265	A	17001	1	777	MSGENVTKVSTFILVGLPTAPG
		ŀ				LQYLLFLLFLLTYLFVLVENLAI
					ļ	ILIVWSSTSLHRPMYYFLSSMSF
	:					LEIWYVSDITPKMLEGFLLQQK
	1	ľ				RISFVGCMTQLYFFSSLVCTEC
						VLLASMGYDRYVAICHPLRY\N
						LACTDFSTAELV/SFILAFIILVFP
	1					LLATILSYWHITLAVLRIPSATG
				1		CWRAFSTCASHLTVVTVFYTA
	1	}				LLFMYVRPQAIDSQSSNKLISA
	1					VYTVVTPIINPLIYCLRNKEFKD
						ALKKALGLGQTSH
16898	47266	Α	17002	2	876	
16899	47267	Α	17003	I	894	
16900	47268	Α	17004	377	501	
16901	47269	Α	17005	3	823	
16902	47270	Α	17006	1	874	,
16903	47271	С	17007	156	287	
16904	47272	A	17008	1	1140	
16905	47273	Α	17009	1299	3135	ISGSVCILLWLQLLPAYNLIIHT
		1				WEHRHSPPGT/RMSRLAKRNA
						WEADR/YAEAKLAASRRGQPSS
						LMSSREGGWGEGSQSRNSSHG
						ALKEAQFIITNAEKSLKDLMEL
						KTMAQELRDECTSLSSRFDQLE
						EKVSVMEDEINEMKQEEKVRE
Ì		1				KRIKRNEQSLQEIWDYVKRPNL
						HLIGVPESDRENGTKLENTLQDI
		İ			1	IQENFPNLARQANIQIQEIQRMP
		1				QRYSSRRATPRHIIVRFTKVEM
	1					KEKMLRAAREKEIQTTIREYYK
						HLYTNKLENLEEMDKFLDTYS
	}					LPRLNQEEVESLKRPITGSEIGAI
						INSLPTKKSPGPDGFTAEFYQRY
						KKELVPFLLKLFQSTEKERILPN
						SFYEASIILIPKPGRDTTKKENFR
						PISLMNINAKILNKILANQIQQHI
						EKLIHHDQVGFIPGMQGWFNIR
						KSINVIQHINRTNEKNHMIISIDA
						EKAFDKIQQPFMLKTLNKLGID
						GTYLKIVRAIYDKLTANIILNGQ
						KLEAFPLKTGTRQGCPLSPLLF
						NIVLEVLARAIRQEKEIKGIQLG
-						KEQVKLSLFADDMIVYLENPIV
						SAQNLLKLISNFSKVSGYKINV
						QKSQAFLYTNNRQTESQIMSEL
16006	47074		17010	190	457	PFTIASKRIK
16906	47274	C	17010	189	457	
16907 16908	47275 47276	A A	17011 17012	107	2600 1493	
10900	17/2/0	I^.	17012	1107	1773	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
ŀ				sequence	}	
16909	47277	A	17013	3	2209	IHTKTPSVCHHHQRPKVDKTTK
10909	4/2//	A	17013	3	2209	MGKKQSRNTGNSKNQSTSPPP
ŀ						KECSSAPAMEQSWMENDFDEL
						_
						REEGFRRSNFSELTEEVRTHRK
						EAKNLEKKIRQM\VTRITNVEK
						SLNDLMELKNMARELRDKCTS
						FSSRFDQLEERVSVIEDQMNEM
						KQEEKFREKRVK\RNEQSLQEIR
						DYVKRP\NLRLIGIPESDEENGT
						KLENTLQDII\QENFPNLARQAN
						IQIKEIQRMPQRYSSRRATPRQI\
						IVRFTEVEMREK\MLRAAREKG
						RVTHKGKTIRLTVDLSAETIQA
-						RRESTRQKVNKDIQELNSAQHQ
					[ADLTDNYRTLHPKSTEYTLFSA
		1				PHHTYSKIDHIVGSKALLSKCK
		1				RTEIITNCLSDHSAIKLELRIKKL
		i				TQNLSTTWKLNNLLLNDYWVH
		ŀ				NEMKAEIKMLFETNENKDTTY
]		QNLWDTFKAVCRGKYIALNAH
			!	}		KRKQERSKIDTLTSQLKELEKQ
			İ			EQSHSKASRRQEITKIRAELKEI
		1				ETQKTLQKINESRDWFFEKINKI
		l				DRQLARLIKKKREKNQIDTIKN
						VLEFLAKAIRQDKEIKGIQLGKE
		l				EVKLFLFAEDMIVYLENPIVSA
						QNLLKLISNFSKVSGYKINVQK
						SQAFLYTNNRQTESQIMSELPFT
						IASKRIKYLGIQLTRDMKDLFK
		l				ENYKPLLNKIKEDTKKWKNIPC
1		١.				SWVGRINIMKMAILPKVIYRFN
		l				AIPIKLPMTFFTELEKTTLKFTW
16910	47278	Α	17014	292	699	
16911	47279	Α	17015	269	803	ALGPCSEALPTWVNENGEEMA
		1				QEIDLSALKELEREAILQVLYRD
		l				QAVQNTEEERTRAAGN*KHTC
						SISGGKGPKNTDWEHKEKCCA
						RCQQVLGFLLHRGAVCRGCSH
						RVCAQCRVFLRGTHAWKCTVC
						FEDRNVKIKTGEWFYEERAKKF
						PTGDNSV\RTSCTGKHETVGGQ
		l				LLQSYQK
16912	47280	A	17016	1	1173	\$7.5 \(\frac{1}{2}\)
16913	47281	A	17017	349	718	AGPEGTTTAECP/I/CQQQRPILS
						LRYGTISWG/DQSATWWQVDY
						IRTLLSWKWQSASAKTTIHGLT
						KCLIHHDIPHSIASD*GTCFMAK
						EVWQWYCFSHSQDSRVQESRG
						GIGSCTTHHHPCSFPN
	<u> </u>	L	<u> </u>	1	1	010001111111111111111111111111111111111

SEQ ID	SEQ ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
16914	47282	A	17018	196	577	GSSGQSESAQPQGLWVLPSRTS
10714	77202	^	17010	170		LAILSTDVNV*QPCFPAGAHQG
						QPLAWPVPFLAWLSWQQKRC*
						KQQRSSRNLVDHRNPGWLHEE
						GDKTPFFHHPFPQLGQSLSQQS
						LLPAGRR*ARGPQQLPSPP
16915	47283	A	17019	1	2523	ELIMOIDE MICH QUEISIT
16916	47284	A	17020	236	558	
16917	47285	A	17021	1610	2004	VTWSAQQMPWAIKNCPTHSPL
	1,203	ļ'``	1,,,,	1010		GQGCLPQPKM*VVQNDFQ*G*
			:			WTHCRKGIKGGQRVGVWPWQ
						TONRGKGTFLTPLHSAVADAM
						LYAL*RGSDLDAAQTLPTAAQ
	į		<u>.</u>			DALYAVVPGGEPTGQLQVRHN
	1					EFALGQ
16918	47286	A	17022	1	735	
16919	47287	Α	17023	88	1871	
16920	47288	Α	17024	1	1299	
16921	47289	Α	17025	2	260	
16922	47290	Α	17026	144	1737	1 1
16923	47291	Α	17027	1	1182	
16924	47292	Α	17028	262	1132	
16925	47293	A	17029	1	484	
16926	47294	Α	17030	3	382	LKGAPDLELDGESRRTWRERK
1		İ				GRKRKERALGRWQV*RSRCA*
						P/STPAALNAPLQGASHSPFRLR
	İ					NCWKGRSVRASSLLRQLAKGG
						CAARRLSWNLCQIVQMYEGHS
		1	:			QQPGLQPRLQRWSSQTLRTA
16927	47295	Α	17031	1	2022	
16928	47296	Α	17032	209	403	
16929	47297	A	17033	213	1993	
16930	47298	A	17034	1	378	
16931	47299	Α	17035	64	1180	SASHSGAPATSLYERGVCPPAL
		1				RRPPAVLVCRAAASCTLLAMS
						DLQEEGKNAINSPMSPALVDVH
						PEDTQLEENEERTMIDPTSKEDP
						KFKELGKVLLDWINDVLAEERI
						IVKQ\LEEDLYDGQVLQKLF/EK
						NLAGLQSLNVAEVTQSEIGQKQ
						KLQKVLEQLH\DLLRPRSWALR
						\WLGNSIHGKNLVGNPQLLVSL
						AMHFRAPIRLPEHVTVQVVVV
	1	1				RKREGLLHSSHISEELTTTEM
						MMGRFERDAFDTLFDHAPDKL
	:					SVVKKSLITFVNKHLNKLNLEV
İ				1		TELETQFADGVYLVLLMGLLE
						DYFVPLHHFYLTPESFDQKVHN
		1				VSFAFELMLGRRPSRNPRLVLK
						TWLTWDLKSPLRVLYNLFTKY
16932	47300	A	17036	3	764	
L10732	1,,200	14.3	1.,000	<u></u>	1,3,	<u> </u>

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
16933	47301	Α	17037	1	1230	
16934	47302	A	17038	1	313	
16935	47303	Α	17039	145	263	IFLLYLKNKVQNK*EIKQHFLG KIMSRDNNTYLTWGTEN
16936	47304	A	17040	5	982	PTFSRAVATMFSRAGVAGLSA WTLQPQWIQVRNMATLKDITR RLKSIKNIQKITKSMKMVAAAK YARAERELKPARIYGLGSLALY EKADIKGPEDKKKHLLIGVSSD RGLCGAIHSSIAKQMKSEVATL TAAGKEVMLVGIGDKIRGILYR THSDQFLVAFKEVGRKPPTFGD ASVIALELLNSGYEFDEGSIIFN KFRSVISYRTEEKP\IFSLNT\VAS ADQHGVSMTDIDADVAAKITQ EY/NIWANIILLTLWKESTTSGG RSARG*QPWDNGQARIAFWRM DLDNLDIWTF\NR\TRQAVITKE
						LIEIISGAASSVKKENSAS
16937	47305	Α	17041	1334	1790	
16938	47306	Α	17042	1	930	
16939	47307	A	17043	2	477	SGLGRLPGPWQEAGSSRGPSSG DMAGVKALVALSFSGAIGLTF/ LHMLGCALEDYGVYWPLFVLI F\HAISSIPHFIAKRVTYDSDAT\S SACRELAYFFTTGIVVSCLWISP VILARVALIK\WGACGL\VLA\G NAVIFPYNSRGFSLYLGRGDDF SW\EQW
16940	47308	A	17044	130	434	
16941	47309	Α	17045	1	1365	
16942	47310	Α	17046	656	839	
16943	47311	В	17047	16	149	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
ĺ	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
16944	47312	Α	17048	1	1317	MTLNEHAAFKHLFNKAHLAPP
		l				LIHLTLSGHSTCFREHRVGVICS
			ŀ			HFAQDLWPEQGREDSFQKVILR
						RYEKCGHENLQLKIGCTNVDES
						LTYKRIHTGEKPCKCEECDKAF
						SKFSILTKHKVIHTGEKHYKCE
						ECGKAFTRSSSLIEHKRSHAGE
						KPYKCEECGKAFSKASTLTAHK
ł						TIHAGEKPYKCEECGKAFNRSS
	;	i				NLMEHKRIHTGEKPCKCEECG
ĺ					İ	KAFGNFSTLTKHKVIHTGEKPY
ĺ						KCEECGKAFSWPSSLTEHKRIH
ĺ		ŀ				AGDKPYKCEECGKTFKWSSTL
						TKHKIIHTGEKPYKCEECGKAF
						TTFSSLTKHKVIHTGEKHYKCE
						ECGKVFSWSSSLTTHKAIHAGE
						KLYKCEECGKAFKWSSRLSEH
						KRIHTGEKPYKCEECGKAFSW
		1				VSVLNKHKKIHAGKKFYKCEE
						CGKDFNQSSHLTTHKRIHTG\E
		1				KPYKCEECGKAFSKASTLTAHK
		1				TIHAGEKPYKCEECGKAFNRSS
		1				NLMEHKRIHTGEKPCKCEECG
						KAFGNFSTLTKHKVIHTGEKPY
		1				KCEECGKAFSWPSSLTEHKRIH
		l				AGDKPYKCEECGKTFKWSSTL
						TKHKIIHTGEKPYKCEECGKAF
		1				TTFSSLTKHKVIHTGEKHYKCE
		1				ECGKVFSWSSSLTTHKAIHAGE
1						KLYKCEECGKAFKWSSRLSEH
						KRIHTGEKPYKCEECGKAFSW
						VSVLNKHKKIHAGKKFYKCEE
16945	47313	Α	17049	2	2183	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
16946	47314	A	17050	3	1440	IKVLLGENRVPNTMAVWLAQ WLGKGASLTSFPPCPTASLLRR
						LGEHIQQFQESSAQGLGLSLGP
						GAAALPKVGWLEQLLDPFNVS
						DRRSHL\QRYWVNDQHWVGQ
						DGPIFLHLHLTSPPPTHTLFTGH
,						PAALAPAWGALVISLEHRFYGL
						SIPAGGRLALSRLFNISSSSPWIC
İ						FGGSYAGSLPSHHAPTF\PRAQR
						IS*HFRIPSSAVRPQVVSRSLMS
						TAIGGSLE/CRRWGLVRGGLGG
						KRGLGCQGRGGPA*VLADIASC
						FPQCRAAVSVAFAEVERRLRSG
						GAAQAALRTELSACGPLGRAE
						NQAELLGALQALVGGVVQYD
						GQTGAPLSVRQLCGLLLGGGG
						NRRGGIYVLCMYIVP*IVLHSLG
						QKCLSFSRAETVAQLRSTEPQL
						SGVGEAPIYLAPSYPFFPGDTDP
:						WHVLSVTQALGSSESTLLIRTG
						SHCLDMAPERPSDSPSLRLGRQ
16015	1.501.5		15051		0.700	NIFQQLQTWLKLAKESQIKGEV
16947 16948	47315 47316	B A	17051 17052	874	2709 1024	
16949	47317	A	17052	1	1602	MDSLWGPGAGSHPFGVHNSRL
10717	',51'	ľ`	1,,033		1.002	SPDLCPGKIVLRALKESGAGMP
						EQDKDPRVQENPGDQRRVPEV
						TGDAPSAFRPLRDNGGLSPFVP
						GPGPLQTDLHAQRSEIRYDQSS
						OTSWTSSCTNRNAISSSYSSTGG
						LPGLKRRRGPASSHCQLTLSSS
						KTVSEDRPQAVSSGHTQCEKV
						AEKAPGQTLALRNDSSRSEASR
						PSTRKFPLLPHRRGEPLMLPPPV
						ELGYRVTAEDLDWEKEAAFQC
						IKSALQVEDKAISDCRPSRPSHT
						LSSLATGAS\DSAQVTSLIPAPFP
					:	AASMDAGMRRTRPGTSAPAAA
]						AAAPPPSTLNRTLGSLLEWMEA
						LHISGPQPQLQQVPRGQNQRSQ
						TSRTSSCPK/LKCHLELLQLYGR
						PPGTKAEEAPASSHCQLTLSSSN
						TVSEDGPEAVSSGHTQCEKTAD
						TAPGQTLGPRGGSPRSQSSRPR
						RHKFPLLPRRRGEPLMLPPPLEL
						GYWVTAEDLDREKEAAFQRIN
						SALQVEDKAISDCRPSRPSHTLS
						SLATGASGLPAISKAPSMDAQQ
						ERHKS
16950	47318	Α	17054	1	1155	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
16951	47319	A	17055	1	1237	 MDSLWGPGAGSHPFGVHNTRL
10931	4/319	^	17033	["	1237	SPDLCPGKIVLRALKESGAGMP
		1				EQDKDPRVQENPGDQRRVPEV
İ						TGDAPSAFRPLRDNRGLSPFVP
						GPGPLQTDLHAQRSEIRYNQTS
						OISWTSSCTNRNAISSSYSSMGG
						LPGLKRRRGPASSHCQLTLSSS
						KTVSEDRPQAVSSGHTQCEKA
						ADIAPGQTLALRNDSSTSEASRP
						STHKFPLLPRRRGEPLML\PPSL
						EVGVPGSLVKELDREKEAAF\Q
						RINSALQVEDKAISDCRPSWPS
						HTLSSLATGTSGLPAISKAPSMD
						AQQETHKSQDCLGLLAPLASA
				1		AGVPSTAPMSGKKHRPPGPLFS
				•		SSDPLPATSSHSQDSAQVTSLIP
						APFPAASMDVGMRRTRRGEPL
						MLPPPLELGYRVTAEDLDQEKE
						AAFQRIKSALQVEDKAI
16952	47320	A	17056	132	835	
16953	47321	A	17057	1	2947	MPEQDKDPRVQENPDDQRRVP
						KVTGDARSAFRPLRDNGVLSPF
						VPRPGPLQTYLHAQRSEIRYNQ
1						TSQNTWTSSCTNQNAISSSYSS
						VGGLLGLKWRRGPAGQESGAG
						MPEQDK\DPRV\QQNPDDHRTV
						PEVTGDARSAFWPLRDNGGLSP
						FVPRPGPLQTDLHAQSSEIRYN
						QTSQTSWTSSSTKRNAISSSYSS
		İ	į			TGGLPGLKQRRGPASSRCQLTL
			ŀ			SYSKTVSEDRPQAVSSGHTRCE
						KAADTAPGQTLAPR
16954	47322	С	17058	229	624	
16955	47323	Α	17059	1	1011	MVSTPATLPSLPKPALMASWG
	:		ļ			VPYDQLTKEEKTRVWFTDGSA
			ŀ			RYAGTTQKWTAVALQPLSRTS
		1				LKDSSEGKSSQWAELQAVYLV
		1				VHFAWKEKWPDVGLYTDSWA
						VANGLAGWSETWEKQDWKIG
						DKEIWGRGMWMDLSEWSKAV
						KIFVSHVSAHQRVTSAEEEFNN
		1				QVDRPL/PVFTQWAHEQSGHSG
						RDGGYSWAQQT\GLPFTKADL
						AMATAECPICQQQRPTLSPLYS
						TIPQGDQPATWWQIDYIGPLPS
ŀ						WKGQKFVLTVIDTYSRYRFAY
						PAHNASAKTTIHGLIECLIHCYG
						IPHSIASDQSIHF/TTKEVQ*WAH
L						AHGIHWSYHVSHHPEAAGL
16956	47324	В	17060	10	597	
16957	47325	В	17061	6	297	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	:			Sequence		
16958	47326	В	17062	1	474	
16959	47327	Α	17063	1	2712	
16960	47328	В	17064	1	738	
16961	47329	Α	17065	3	772	DPADPMVLEVSEADRDA\VPIS
						ESQQRPLGFWSKALPSSANNYS
						FFKRQLLACYWVLVEIEHLTM
						GHQVTMRPELPIINCVLSDPCSH
						KVGHAQQHSIIKWRWYIHYQA
						QAGPEGTSNLHEEVAQMPTVS
						TSATLPSLPEPTQMAL\WEVPY
						D\QFTEEEKTGAWFTDGSAHYT
						GTT*KWTVAALQPLSRTTLKDS
						CEGKSPHHPVIAQWAHEQSGH
						GGRDGGYLWAQQHGFPLTKA
		1				DLAMATAECPICQQQKRTLNP
16962	47330	Α	17067	1667	2012	LSPHWPPGLPPSALIYAFLGRSS
ļ						RNKRRLMKTMKVEACEDLSAP
						KYLRIQLPTGSPRRRSEAPEVCI
						FQGSS\FSLVHPPPLLSYPSARR*
						PQTPPRPPPRHPSLHPLHPPSAQ
16963	47331	Α	17068	5	1074	RHSALGTGGSSSCRQQRPRDT
	İ					WVPETTILTAWSS/RLQQLGPW
						LQVTVAGQVNTLAGIQKTIHAE
				1		HCNVTE\RPPTGHEALAPPCEQ
		1				GPPQVLDSNAVAPTADCGYQL
ļ						STAINRNQELCGRKPAVSPSR/G
						ISPGSPILRPAVHPGHYCLQQRL
						SGQKSNRRTTQKSFQKLHVDK
						RKPENFSKESDHLLIYSREGNPD
						VASLATPPGIWMMSATDLFYG
						DPFAFPGGGPFTLSLERGPSVIL
						GTDASEELLFTKASLAVSTSVF
						QFSTGKIYTEWVWQKNKSFLSP
						HFPTKPMASPVKGKHSTSLLQR
						LNYRDRWQCLQEGAGLALLSA
						HMKVPTGSKEPQQAGKLPSKP
						RKQTTPTFKD
16964	47332	Α	17069	2	699	CSACPVSHPWLLPWQSGSAFSL
						P*RALQRPPQHGHRLSKPMQSP
						RLSVLLRPWPRRPLPTDLLY*PS
		1				TRGHIHLHDRHHGAPSVSRRW
						GDKASPPQFPRYSRDSK/RRPPQ
		1				HGEVGTVFTNNKGGQS/PATST
						WEVQSAECQKAPEAGPTPPSTA
		1				GDAEGANALRQESEGNWGSEA
		1				KSPHVAEAGSWRPQQGSCHHL
						HPPDKTKDAARKTHSPGDNGC
						QPRPPRPCKTLWSLSRI
16965	47333	Α	17070	223	1611	-
16966	47334	A	17071	12	310	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
16967	47335	A	17072	31	430	LQAVRLFTLPTTSSGL*KLRTVP
						TWPSPPSCFPHFQSGPRTTRAPT
		İ				PSTI\PGYSGSYSSGPPSLCPRTS
						RRAPPRHRVPAGRERLHIRVPP
						YPRPERHLLRVSRRGILGNNRV
						GRLPESRAIAGRKEEKTDTK
16968	47336	A	17073	1	474	
16969	47337	A	17074	153	1224	RVFSESVCSPVRNLEFLWRFAF
10909	4/33/	Ι^	17074	133	1224	PLAPAGRCPPGVPLQTSPRDTD
						AHRSSPLPPARASPGQVAAAYR
	i					1
	ŀ	ŀ				WARCPGCGGRKPRSSGSWQLC
						RCPTLPPPPRGSRSSGRC/RTWP
				}		SPPSCFPHFQSGPRTTRAPTPSTI\
						PGYSGSYSSGPGR*GLSPLHAA/
						VSPPLPPGGP*GSWARAGLGSI
		1				ASAHSPCPLCRSLIRSRS*QTCT
						RSPT*NCEVPPSAP*AASPLRTM
		ł				FALVRTAGLKVHLLPLGYCTT
						MS*SSSMPQTVPVVVKVSNIPS
		l				VHPP*PCCKDCTISRSRSIFTRSPI
		1				CNPPGFLLPFCSPSTGQ*SL*KEP
						PLASWTHFRSDVLLLFSVSMNG
		1				STLSLGCPSQKAVIALVQVT
16970	47338	Α	17075	215	1246	
16971	47339	Α	17076	566	675	QPCRQQ/CLQPGSPPRFHARTLT
						VTLVHTRTHTHAHAH
16972	47340	Α	17077	184	584	
16973	47341	Α	17078	1	1167	
16974	47342	Α	17079	849	2266	
16975	47343	Α	17080	1505	1891	DAEKAFDEIQQ/PFMIKILSRISI
	1	1		1		QGTYLNVIKTIYDKPTANTILN
		1				VEKLKTFPLRTGTRQGCPLSAL
	ı	1				LFNMVLEVIARAIRQEKGIKGIQ
]	1	1				ISKKEVKLSLFADDMIVHIENPK
l		1				DSSRKLLELIKEFSKV
16976	47344	A	17081	1	842	
16977	47345	A	17082	155	531	GNLWSVDLRPGTPLRQNFRGTI
1	1,,3,5	` `	17002	1.55		RQQHSRFTKNHCSQTPLLIPRQ
						TGSGVDLSKLQQTCS*GSCLVC
						TIDLANAFFSIPVHKA/HQKQFA
						1
						FSWQ\YTFTVLPRLTWLQPC*V PNLPAAETNTEPSNGT
16978	47346	A	17083	1	1773	PNLPAREINIEPSNGI
16979	47347	В	17084	598	1428	
					227	
16980	47348	A	17085	3	<u> </u>	
16981	47349	Α_	17086	1	207	<u> </u>

SEQ ID	SEQ ID NO:	1	SEQ ID NO:	í .	1	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
		<u> </u>				
16982	47350	Α	17087	1	789	RDLQPFTSVTVHCRKGNDQTF
	1					GGPLDAGSELTLIPGDPKHHCG
						PPVKVGAYGGQVINGVLA\HPL
						IWLVQKTDGS/WRMTVDYCKL
		1	-			NQVVIPIAAAVSDVVSLLEQINT
		1				SPGTWYAAIDLANAFFSIPVHK
						AQQKQFAFSWQGQQYTFTVLP
						QWYINSPALCHNLIRRDLDCFS
		1				LPLDITLVHYIDDIMLIGSSEQE
						VANTLDLFVRHLRARGWEINPT
		1			-	KIQGPSTSVKFLGFQWCGACQA
						IPSKMRDKLLHLVPPTTKKEAQ
16983	47351	Α	17088	1	435	
16984	47352	A	17089	2	808	PLIWLVQKTDGS/WRMTVDYC
		1				KLNQVVIPIAAAVSDVVSLLEQI
		ļ				NTSPGTWYAAIDLANAFFSIPV
		ŀ				HKAQQKQFAFSWQGQQYTFTV
						LPQWYINSPALCHNLIRRDLDC
						FSLPLDITLVHYIDDIMLIGSSEQ
						EVANTLDLFVRHLRARGWEINF
						TKIQGPSTSVKFLGFQWCGACQ
						AIPSKMRDKLLHLVPPTTKKEA
						QCL\QLLACY/WALVETEHLTIS
						HQVTMRPELPIMNWVLFDPSSH
						KVGCAQQHSIIKWKWYVHDW
						ARAGPEGT
16985	47353	Α	17090	2	544	
16986	47354	Α	17091	1	633	MTVDYRKFNQVVTPMAA/AVP
						DAVSLLEQINTFPGTWYAAIDL
						ANAFFSIPVHEAHQKQFAF\LPQ
						GYINFPALCHNLIRRELDFFLLL
		İ				QDITLVHYIDDILLIGSSEQEVV
						NTLDLLIHKRSKEAEHTAASRIR
						VSCLPEQKSHEQTLPWEQVPSS
						GDIKEYFPNAFVLLTTASLQGQ
						DNTSQLQLTWKAPEDIKMSKT
						DADADEEIEALRG
16987	47355	Α	17092	1	3228	MVTSSLTVAITSLVVVITSLVVII
]]	TSFVVVTTSLVVVITSLVVVITSI
						AIAIATLVVVLTFTVVVLTSLV
1			1	•		VVTTSLVMVITPPFAVITSLGVV
						ITSLVVIITCLVLVITALGVVITS
						LGVVVTSLVMVITPLVAVITPL
						GVVITSLVVVITSLVMVITSLTV
						VITSSIVVITPLVMVITSLTVIITS
					1	LFVVITPLVMVITFLIVVNTSLV
						VVLITLIVSPPPSLCSLVVVITFL
						VVAITSLVVVITSLVVVITSL
16988	47356	A	17093	1	471	
16989	47357	A	17094	1	1590	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
16990	47358	A	17095	332	1704	RGYVFCSWKKTDGS/WRMTVD YCKLNQVVIPIAAAVSDVVSLL EQINTSPGTWYAAIDLANAFFSI PVHKAQQKQFAFSWQGQQYTF TVLPQWYINSPALCHNLIRRDL DCFSLPLDITLVHYIDDIMLIG\P RQLLACY/WALVETEHLTISHQ VTMRPELPIMNWVLFDPSSHKV GCAQQHSIIKWKWYVHDWAR AGPEGTTTPVISQWPHEQCGHG GRDGGYAWAQQCRLPLTKAD LNTATAKRPICQQQRPTLSPQY GTIPQGDQPATWWWVDYMGS LPSWKGQRFVLTGIDTYSGYGF AYPACNASAKTAICGLTECLIH HHDIPHSIASDQGTHFMAKEVR QWAHDHGIHWSYHVSHHPEA AGLIEWWNGLLKSQLQCQLGD NTWQGWGKVLQKVVYALNQH PIYGTVSPIAKIHRRVADSLKGG SGSWVWGFSTLAALDNDLGTF RKVRAPSTN
16991	47359	В	17096	1	2061	
16992	47360	Α	17097	1	1623	
16993	47361	Α	17098	1	741	
16994	47362	Α	17099	213	679	
16995	47363	A	17100	33	853	
16996	47364	A	17101	3	1249	
16997	47365	A	17102	17	770	
16998	47366	Α	17103	130	469	
16999	47367	Α	17104	1	1425	
17000	47368	Α	17105	1	882	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
17001	1472.60	1 4	17106	227	1519	VTLIKMNAMLETPTLS\AVSDG
17001	47369	Α	17106	227	1319	· ·
						VKLSAVAAVLYVIVRCMNLKS
						ATAAPDLYLQDSGLSRFLLKSC
						PLLTKEYIPPLIWGKSGHIQTAL
ļ		l				YGKMGRVRSPHPYGHRKFITM
		:				SDGATSTFDLFEPLAEHCVGDD
						ITMVICPGIANHSEKQYIRTFVD
	E					YAQKNGYRCAVLNHLGALPNI
						ELTSPRMFTYGCTWEFGAMVN
						YIKKTYPLTQLVVVGFSLGGNI
						VCKYLGETQANQEKVLCCVSV
		1				CQGYSALRAQETFMQWDQCRR FYNFLMADNMKKIILSHRQALF
						`
						GDHVKKPQSLEDTDLSRLYTAT
						SLMQIDDNVMRKFHGYNSLKE
						YYEEESCMRYLHRIYVPLMLV NAADDPLVHESLLTIPKSLSEKR
					1	ENVMFVLPLHGGHLGFFEGSVL
			}			FPEPLTWMDKLVVEYANAICQ
17000	47270	 	17107	21	205	WERNKLQCSDTEQVEADLE
17002	47370	Α	17107	21	385	SRSSSVVRQEPGRADGQRRRRR
1						GGPGLRSPGERQQPRPARPCRR
						AHHPGDPAGAQ/PPPKTGKTCG
						APKATTRSTRCSISR*PPFPRSSC
						PTSPRATPSSSASSTAAMRRWL VTCRMPNWRA
17003	47371	A	17108	158	423	VICKWIFNWKA
17003	47371	A	17109	1	1851	
17004	47373	A	17110	1	1104	
17006	47374	A	17111	66	432	
17007	47375	A	17112	103	3531	
17008	47376	A	17113	1	1311	
17009	47377	A	17114	1	750	
17010	47378	A	17115	183	373	QPQATTLVHPVHVA*LF*VHW
		1				QNRYRAPDNQGYRHWRCLLPA
						LPDAGR/VRLSASC*KPPNSRG
17011	47379	Α	17116	344	480	SPKDQEPSISKKV***TSLPTSSR
						SLCLPPARMHFWVLAALLSFA
17012	47380	Α	17117	3	3565	HEARRGLKMAACGRVRRMFRL
						SAALHLLLLFAAGA\RNSPARA
		1				SHSQGQGPGANFVSFVGQAGG
						GGPAGQQLPQLPQSSQLQQQQ
						QQQQQQQQPQPPQPPFPAGGPP
						ARRGGAGAGGGWKLAEEESCR
						EDVTRVCPKHTWSNNLAVLEC
		1				LQDVREPENEISSDCNHLLWNY
		1				KLNLTTDPKFESVAREVCKSTIT
		1				EIKECADEPVGKGYMVSCLVD
						HRGNITEYQCHQYITKMTAIIFS
		1				DYRLICGFMDDCKNDIN
	<u> </u>			<u></u>	<u> </u>	

=Unknown, nucleotide otide insertion) *SGPAPRL/C GVCL*RHW RLSSR SPETAQPPP NLLNMAKL FLDSDYPPL
*SGPAPRL/C GVCL*RHW RLSSR SPETAQPPPP NLLNMAKL
GVCL*RHW RLSSR SPETAQPPPP NLLNMAKL
GVCL*RHW RLSSR SPETAQPPPP NLLNMAKL
GVCL*RHW RLSSR SPETAQPPPP NLLNMAKL
RLSSR SPETAQPPPP NLLNMAKL
SPETAQPPPP NLLNMAKL
NLLNMAKL
IGLKVRKSF
VEKLYPEAE
EFYEYHALM
QMRWWPE
PPETAQPPPP
NLLNMAKL
TLDSDYPPL
HGLKVRKSF
VEKLYPEAE
TPLGRARAW
YLRCLIIQR
MEEEGAVI

DNPSVSNRL
ICSD*SQVL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
17034	47402	Α	17139	476	2934	TEPKTKTT*LSQ*MQKKPLTKF
		1				NNPSC*KLSIN/IVLEVLARAIRQ
						EKEIKGIQLGKEEVKLSLFADD
	-					MIVYLENPIVSAQNLLKLISNFS
						KVSGYKINVQKSQAFLYTNNR
						QTESQIMSELPFTIASKRIKYLGI
						QLTRDVKDLFKENYKPLLKEIK
						EDTNKWKNIPCSWVGRINIVK
						MAILPKVIYRFNAIPIKLPMTFF
						TELEKTTLRFIWNQKRARIAKAI
						LSQKNKAVGITLPDFKLYYKAT
						VTKTAWYWYQNRDIDQWNRT
		1				EPSEITPLIYNYQIFDKPEKNKQ
		1				WGKDSLFNKWCWDNWLAICR
						KLKLDPFLTPYTKINSRWIKAL
						NLRPKTIKTLEENLGITIQDIGTG
			ļ			KDFMTKTPKAMATKDKIDKW
						DLIKLKSFCTAKETTIRVNRLPT
						KWEKIFATYSSDKGLISRIYNEL
						KQIYKKKTNNPIKKWEKDMNR
						HFSKDDILAAKKHMKKCSSLAI
						REMQIKTTMRYHLTPVRMAIIK
						KPGNNSQKDIPDWRNGPKRKD
						RQITTFRGPTKLQVVVPTLGAA
1						PGRGCPWPRRPRCPGREGEELH
						PALGGERGQRQAAGSGGARRL
						APSAGDGLCVGPVEDTGETQT
						WKTGQPGLREPWWGWRPTEPP
						PAPSSHSEEWPRRPPVRRAPRP
						AGEHAGEGGAAVADTTAFSPW
						LPRVGGWHLQVHRGGRSCAGA
						RRFGGSRGWDLGDLSSAPVASL
17035	47403	Α	17140	1	2298	
17036	47404	Α	17141	3	482	
17037	47405	A	17142	1	1026	,
17038	47406	Α	17143	1	623	
17039	47407	Α	17144	25	392	ESLKGADPKFLRNMRFANKHN
					Í	KKGLKKMQANSARATSARAD
						AIKALVKPKEGKPDIPKGVSRK
						LDPLAYIAHP\RLGKCAHSCIAK
						GLKLCRLQAKDKDQTKAQAA
						APA*APKGARTPTKASE
17040	47408	Α	17145	1	507	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
17041	47409	Α	17146	2	811	YRFSGPTIRSIQNHTTHNQGRK
						WHKKGIKKPRAQRYESLKGPE
						V*PSGQGVTCLLRADWLSPLLG
						SGTY\VGPKFLRNTRFAKKHNK
						KGLKKMQANNAKAMSAR\AE
						A\IKALVKPKEVKPKIPKGG\SR
						KLRSTLPDIVHP\KLGKSARAPY
						LPRGFRLCLPNANAKSISSQGQ
						GQGQGQGSNQGPGFKPQLQVP
						ASGVPNVPRALTKGFQSRYSLA
						QHEGQERTWVRPPTPRPWGYH
	1					FAWGWGSPVAICTNKPEAGKR
						RNEKEKLCQGDGY
17042	47410	Α	17147	1	636	
17043	47411	Α	17148	169	410	
17044	47412	Α	17149	385	1359	QTNSPNCVRGRVSPPVLPGSSS
						MDVLASYSIFQELQLVHDTGYF
1						SALPSLEETWQQTCLELERYLQ
				1		TEPRRISETF\GEDLD\CFL\HAFP
		1				SPCIEESFRRLDPLLLPVEAAICE
						KSSAVDILLSRDKLLSETCLSLQ
						PASSSLDSYTAVNQAQLNAVTS
	1					LTPPSSPELSRHLVKTSQTLSAV
		1				DGTVTLKLVAKKAALSSVKVG
						GVATAAAAVTAAGAVKSGQS
						DSDQGGLGAEACPENKKRVHR
						CQFNGCRKVYTKSSHLKAHQR
						THTGEKPYKCSWEGCEWRFAR
						SDELTRHYRKHTGAKPFKCNH
						CDRCFSRSDHLALHMKRHI
17045	47413	Α	17150	1	948	
17046	47414	Α	17151	400	446	PSFLR*PSFWSREFH
17047	47415	Α	17152	7212	10914	ADSGCKL/PSSC*HVLCA/WVTY
						MVMAAILHKLDQLDASRGRAI
						PFCGQRGRFKYEILVKTGWGR
		1				GSGTTAHVGIMLYGVDSRSGH
		1				RHLDGDRAFHRNSLDIFRIATP
		1				HSLGSVWKIRVWHDNKGLSPA
		1				WFLQHVIVRDLQTARSAFFLVN
		1				DWLSVETEANGGLVEKEVLAA
		1				SDAALLRFRRLLVAELQRGFFD
		1				KHIWLSIWDRPPRSRFTRIQRAT
		1				CCVLLICLFLGANAVWYGAVG
		1				DSAYSTGHVSRLSPLSV
17048	47416	В	17153	1	14934	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
17040	47417	 	17154	999	3009	ADSGCKL/PSSC*HVLCA/WVTY
17049	47417	Α	17134	1999	3009	MVMAAILHKLDQLDASRGRAI
						PFCGQRGRFKYEILVKTGWGR
		1		ŀ		GSGTTAHVGIMLYGVDSRSGH
		1				RHLDGDRAFHRNSLDIFRIATP
		1				HSLGSVWKIRVWHDNKGLSPA
	1					WFLQHVIVRDLQTARSAFFLVN
						DWLSVETEANGGLVEKEVLAA
			1			SDAALLRFRRLLVAELQRGFFD
	ļ					KHIWLSIWDRPPRSRFTRIQRAT
						CCVLLICLFLGANAVWYGAVG
						DSAYSTGHVSRLSPLSVDTVAV
						GLVSSVVVYPVYLAILFLFRMS
		ļ				RSKVAGSPSPTPAGQQVLDIDS
						CLDSSVLDSSFLTFSGLHAEVRT
						LLGVLGWAGGPAALAQLGLKT
						LCTSQQAFVGQMKSDLFLDDS
						KSLVCWPSGEGTLSWPDLLSDP
						SIVGSNLRQLARGQAGHGLGPE
						EDGFSLASPYSPAKSFSASDEDL
						IQQVLAEGVSSPAPTQDTHMET
						DLLSSLCAEVHTALPLHCSCLQ
						DSVQWHAVQGLPAFCRGDPVT
						QEAQETARPAAQNLVLIPLHAA
						PHQAVAEIDALYDVYLDVIDK
		l				WGTDDMLFLGDFNADCSYVR
						AQDWAAIRLRSSEVFKWLIPDS
ŀ						ADTTVGNSDCAYDRIVACGAR
						LRRSLKPQSATVHDFQEEFGLD
		ľ				QTQASGPWGGAGLGSPPPGRT
						QAAGRALAISDHFPVEVTLKFH
17050	47418	Α	17155	242	363	TKRQENGRKKQDRTPTEQQLR
						VK*KSRSTKKRRKQSRSES
17051	47419	В	17156	1	12879	
17052	47420	Α	17157	3	596	NLLFPVRSERGPHRPTPLSWLW
						LRPCELNKLAAGGLAAGSSRA
						GALYLLIYIPRRPARAAPLCPAA
		1				ATGAPGASAGSASPASARPPGL
		1				ARAAPGPLRGRRPRSPRTPFCV/
						WLGVSLLLS\YQHYFAEPFESFQ
		1				VCCRHCGILCLTIRVHFPCLSAN
]						PFHFSQLHLGPRPGKACHSVGL
						QIYNSQAHSHPSQQERQLFC
17053	47421	Α	17158	1	235	FFFVLYNLFLFVITSIPLQKVSIL
		1				TQRQLVAEENVKCCTV\PEPQ**
						*LSNYPIFVEMGFCHVGPGWSR
						TPTPKSVSR
L	<u></u>]	L	<u> </u>		TPTPKSVSR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
17054	47422	A	17159	256	4064	ADSGCKL/PSSC*HVLCA/WVTY
						MVMAAILHKLDQLDASRGRAI
				1		PFCGQRGRFKYEILVKTGWGR
]						GSGTTAHVGIMLYGVDSRSGH
						RHLDGDRAFHRNSLDIFRIATP
						HSLGSVWKIRVWHDNKGLSPA
						WFLQHVIVRDLQTARSAFFLVN
į						DWLSVETEANGGLVEKEVLAA
						SDAALLRFRRLLVAELQRGFFD
	1			İ		KHIWLSIWDRPPRSRFTRIQRAT
						CCVLLICLFLGANAVWYGAVG
						DSAYSTGHVSRLSPLSV
17055	47423	A	17160	2	347	LTSRGP/LCCVLLICLFLGANAV
						WYGAVGDSAYSTG/LCVQAEP
						AERRHSRCWPGVQRGCLSRLP
						GHPLSLPDVPEQGGWEPEPHTC
1						RAAGAGRRQLPGLNPCWTAPS
						SRSEASTLR
17056	47424	Α	17161	9415	9994	ADSGCKL/PSSC*HVLCA/WVTY
						MVMAAILHKLDQLDASRGRAI
			:			PFCGQRGRFKYEILVKTGWGR
]		:			GSGTTAHVGIMLYGVDSRSGH
	•					RHLDGDRAFHRNSLDIFRIATP
ŀ		1				HSLGSVWKIRVWHDNKGLSPA
						WFLQHVIVRDLQTARSAFFLVN
						DWLSVETEANGGLVEKEVLAA
						SKASFRVPTPSRSPVALPAPAGG
17057	47425	Α	17162	1030	1222	
17058	47426	Α	17163	2	2762	
17059	47427	A	17164	25	115	
17060	47428	A	17165	1	383	
17061	47429	Α	17166	120	596	IKAPCASFPTSGFPGQRRSFVGR
						SSSTPEECGHSELGRNEDKASR
		1				GRIRQQ/PFAVLKYLLFCSLRW*
		1				YPGKQGLEWTSSKLQQTCRKY
	1					RERHKDTPQEEQLQDT*LSG/V
						TK/G*NEGKNVKGSQRERSGYS
						QREAHQTNSRSLGRNSTSQKRV
				1		GASIQRS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
!	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
17062	47430	Α	17167	1	2236	RPSGPQPLPGSVRHPRPVLRRPL
						PRAQGSSSSFRPRPPFAPDTMD
						KFWWHAAWGLCLVPLSLAQID
		İ				LNITCR\FAGVFHVEKNGRYSIS
		l				RTEAA\NLCKAFNSTLPTMAQM
		1				EKALSIGFETCRYGFIEGHVVIP
		İ				RIHPNSICAANNTGVYILTSNTS
						QYDTYCFNASAPPEEDCTSVTD
		-				LPNAFDGPITITIVNRDGTRYVQ
						KGEYRTNPEDIYPSNPTDDDVS
						SGSSSERSSTSGGYIFYTFSTVH
						PIPDEDSPWITDSTDRIPATSTSS
						NTISAGWEPNEENEDERDRHLS
		ļ				FSGSGIDDDEDFISSTISTTPRAF
		1				DHTKQNQ\DW\TQW\NPSH\SNP
		l				EVLL\Q\TTTRMTDVDRNGTTA
						YEGNWNPEAHPPLIHHEHHEEE
						ETPHSTSTIQATPSSTTEETATQ
	1	ŀ				KEQWFGNRWHEGYRQTPREDS
						HSTTGTAAASAHTSHPMQGRT
						TPSPEDSSWTDFFNPISHPMGRG
						HQAGRRMDMDSSHSTTLQPTA
		İ				NPNTGLVEDLDRTGPLSMTTQ
						QSNSQSFSTSHEGLEEDKDHPT
						TSTLTSSNRNDVTGGRRDPNHS
						EGSTTLLEGYTSHYPHTKESRT
						FIPVTSAKTGSFGVTAVTVGDS
						NSNVNRSLSGDQDTFHPSGGSH
						TTHGSESDGHSHGSQEGGANTT
						SGPIRTPQIPEWLIILASLLALALI
						LAVCIAVNSRRRCGQKKKLVIN
					1	SGNGAVEDRKPSGLNGEASKS
17063	47431	Α	17168	1	1980	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
17064	47432	A	17169	362	2551	EDFSTSATTPSSSPRELPSGQGL
						WLCWLELLPSFTRNGVVHTGP
		1				C*GLC*RWQLSPQPSLPSNFGM
						KISDMATLITTVPAA/FSSSSDW
			[NTPAPTQSPEEVRRLHLCTSFM
						DMLKVGGQGEGAAAGLVQNP
					·	SGHALGCLDSAASGISDPSVAV
						LLENVPNQRGESLRCVPVYLGQ
						WNCSKETGMDSGQFGEEKGAH
						DPSPTYYHACLSTQWWPSPESP
						YPQPSGPQVKEPRNPGKSISML
						CLWIRYHAADSGSSRAKLPLEE
						RLRPFSPTHDCLAFPLPLEKRPE
				1		GNVGSRCPSGTKLPEEGSGSNI
						CCSAIFPVLQPPLVIPRQTRSGV
		1				DLQQTPTDLQLRDLTDRSSLPA
						MEQSWIENDFDELNRSRLQKA
						`
						NKLENLEETDKFLDTYTLPRLN
						QEETESLNRPITGSEIEAIINNVP
						TKKSPGPDGVTAKFYQRYKEE
						LRTVNKNHMIISIDAEKAFDNIQ
						QPFMLKTLNKLGIDATYLKIIRA
						IDDKPTANIILNGQKLEAFPLKT
				1		GTRQGCPLSPLLFNIVLEVLAR
						AIRQEKEIKGIQLGKEEVKLSLF
		1				ADDMIVYLENPIVSAQNLLKLI
						NNFSKVSGYKINVQKSQEFLYN
						NNRQTENQIMSELPFTIASKRIK
						YLGIQLTRDVKDLFKENYNPLL
						NEIKKDTNKWKNIPCSWIGRINI
						MKMAILPKSGPSAARLLEFAGG
						PLQTLFAWVSPEEAAEQQILPN
17065	47433	A	17170	1	483	
17066	47434	A	17171	52	321	·
17067	47435	A	17172	274	393	
17068	47436	Α	17173	304	415	
17069	47437	Α	17174	321	796	
17070	47438	A	17175	1	198	CONTRA A TO LOD COO A TODO DO CO
17071	47439	Α	17176	2	454	CRPVPAATNGRCFPATPRDPGP
			ļ			RMRRACAWGRSKGAGRCLRR
						ATRRDKMAKKESILDLSK\YID\
						KTIRVKFQGGRK\ASGILKGFEP
						NLL\NLVLTVTI*YMRDPDDQY
		İ				KLTED\TRQLG\LVVFRG\TSLVL
						\ICPQDGMEAIP\NPFIQQQDA
17072	47440	Α	17177	1	1806	
17073	47441	Α	17178	1215	2065	
17074	47442	A	17179	217	441	
17075	47443	Α	17180	117	296	DWSDIQKLG*SLSLSKEIFQL*IV
						SPPALHPAKLLHRSA*LSVPQIG
				<u> </u>		RECQGCGWTAMC

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
17076	47444	Λ	17181	2	138	LRTAPEFPGRRFRGCAAAAAES
17070		`	1,101			VAALL*STPPCRSGDSLCGKSR
17077	47445	Α	17182	222	760	
17078	47446	A	17183	192	244	YIQPVF*GWTLGCCQFF
17079	47447	A	17184	1	990	
17080	47448	Α	17185	1	2196	
17081	47449	Α	17186	459	1387	
17082	47450	Α	17187	2	454	
17083	47451	Α	17188	1	957	
17084	47452	Α	17189	398	553	
17085	47453	A	17190	2	293	
17086	47454	В	17191	53	182	
17087	47455	Α	17192	3	1114	
17088 17089 17090 17091	47457 47458 47459	A A A	17194 17195 17196	2 185 3	291 307 392	SVVEFIPSCAVVLWSSVLLSPVR LLSYLSGVPPGHPMEDSMDMD MSPLRPHNYLFGCELKAGQDY HFKVDNDENEHQLSLRTVSLG\ AGAKDELH\IVEAEAMYYEGSP IKV/THLATLKMSVQPT\VSLGG \FEITPPVV\LRLKCGSGPVHISG QHLVAV\EEDAESEDEEEEDVK\ LSLSISGKRSAP\GGGSKVPQKE RKTCMLMKDDDD\DDEEDGYD EDD\DDDDDDF\DDEEAEEKAP SERNLYRDTP\AKNAQKSNQNG KDSKPSSTPRSKGQESF\KKQEK TPKTPKGPSSVEDIKAKMQASI A/EKGGSLPKVEAKFINYVKNC FPMTDQDAIQDLWQWRKSL VRGGQAFRQRPAPPLPVTRSAH RGRRPTRASQPSLFHLLLREGK MSESSSKSSQPLASKQEKDGTE KRGRGRPRKQP\PVSPGTALVG
17092	47460	С	17197	76	378	SQKEPSEVPTPKRPRGRPKGSK/ NKLEKEEEEGISQESSEEEQ
17093	47461	В	17198	1803	1905	
17094	47462	Α	17199	1	416	
17095	47463	Α	17200	30	407	
17096	47464	A	17201	1	993	
17097	47465	A	17202	1	1287	
17098	47466	Α	17203	1	2004	
17099	47467	Α	17204	1	1368	
17100	47468	В	17205	1	3035	
17101	47469	В	17206	1	981	
17102	47470	A	17207	1	2017	
17103	47471	Α	17208	1	1069	
17104	47472	A	17209	1	2463	
17105	47473	Α_	17210	1	324	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
17106	47474	I A	17211	51	1528	GREASKMAQTQGTRRKVCYY
17100	4/4/4	^	17211		1320	YDGDVGNYY*G\QGHPMKP\HR
						IRMTHNLLLNYGLYRKMEIYRP
						HKANADEMTKYHSDDYIKFLR
						SI/RVPDNIVGSTSKQMQRFNVG
						EDFPSIPMACFEFCQLSTGGSVA
						KCL*NFNKQQTDIAVNWAGGL
						HHAKKSEASGFCYVNDIVLAIL
						ELLKYHQRVLYIDIDIHHGDGV EEAFYTTDRVMTVSFHKYGEY
	-		1			FPGTGDLRDIGAGKGKYYAVN
e.	}					YPLRDGIDDESYEAIFKPVMSK
ľ						VMEMFQPSAVVLQCGSDSLSG
		1				DRLGCFNLTIKGHAKCVEFVKS
						FNLPMLMLGGGGYTIRNVARC
						WTYETAVALDTEIPNELPYNDY
		Ì				FEYFGPDFKLHISPSNMTNQNT
						NEYLEKIKQRLFENLRMLPHAP
						GVQMQAIPEDAIPEESGDEDED
						DPDKRISICSSDKRIACEEFSDS
						EEEGEGGRKNSSNFKKAKRVK
						TEDEKEKDPEEKKEVTEEEKTK
17107	17.475	 	17010		400	EEKPEAKGVKEEVKLA
17107 17108	47475 47476	A	17212 17213	1	492 1851	
17108	47477	A	17213	1	753	
171109	47477	B	17214	427	614	
17111	47479	В	17216	1	711	
17112	47480	A	17217	750	1173	
17112	47481	A	17217	1	1224	
17114	47482	A	17219	1242	1743	
17115	47483	A	17219	423	1739	
17116	47484	A	17221	117	1183	, , , , , , , , , , , , , , , , , , ,
17117	47485	A	17222	1	377	FRRGAW/VPLLTRWGFRPRGGC
11/11/	47465		17222	1	377	ISNDSPRRS/IHYRETKPEPPVRV
	İ					CARGLORTPSAGHRAVLRMPR
		1				RGRTRPGAPQGGAALGGPARD
						RECACAGRGAQTFALRKNCVC
		1				KMPSGRGRTCSKNFSYFPS
17118	47486	Ι_Δ	17223	331	1134	KMI SUKUK I COKNI O I FI FO
17118	47487	A A	17224	178	1200	
17119	47488	A	17225	1	367	
17121	47489	A	17226	85	3360	
	47489		17227	8	930	
17122	47491	A	17228	· · · · · · · · · · · · · · · · · · ·	765	
17123		A		1120	509	
17124	47492	В	17229	120 232	3187	
17125	47493	A	17230 17231	1	999	
17126	47494	Α	L	1	325	
17127	47495	A	17232	1	535	
17128	47496	В	17233	271		
17129	47497	Α_	17234	271	634	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
17130	47498	В	17235	301	523	
17131	47499	A	17236	1	1185	
17132	47500	Α	17237	178	497	LPVPAYELNGSQQLPLLLGAAA
						IGYLAYKRFYVKDHRNKAMIN
						LHIQKDNPKIRHAFDME\DLGD
						KAV\YCRCWRSKKFPFCDGAHT
						KHNKE\TGDNVGPLIIKKKET
17133	47501	В	17238	184	395	
17134	47502	Α	17239	1	103	PAWHEGQPQPESECPPQRQCIC
						SHGRGTPARCGVSY*ART/PAG
]					ESP*TQQDLRAS*LGTRANHSQ
						RASARHKDSASAVTGAGHQRD
						AE
17135	47503	Α	17240	447	667	HRPRRRFQIKPQSQVPEHAPLPL
						PTASDPP/AHSPRCPSESRLSPGG
						QRLGFPSSVLNCSKCSWQLAAC
1						MGFSF
17136	47504	Α	17241	265	890	
17137	47505	Α	17242	1	1374	
17138	47506	Α	17243	1	6189	
17139	47507	A	17244	1	8277	
17140	47508	Α	17245	178	1177	
17141	47509	Α	17246	1	447	
17142	47510	Α	17247	1	465	
17143	47511	Α	17248	3	3910	
17144	47512	Α	17249	2	1871	
17145	47513	Α	17250	1	666	
17146	47514	Α	17251	1	635	FPGRRFRRAVWCTHCCAPTPSG
						PVLPHSAAMSFLKSFPPPGPAE
				-		GLLRQQPDTEAVLNGKGLGTG
	:					TLYIAESRLSWLDGSGLGFSLE
						YPTISLHALSRDRSDCLGEHLY
						VMVNAKFEEESKEPVADEEEE
,						DSDDDVEPITEFRFVPSDKSAF*
						FPCSVEAMFTAMCECQALHPD
						PEDEDSDDYDGEEYDVEAHEQ
						GQGDIPTFYTYEEGLS
17147	47515	Α_	17252	1	2796	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217		of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
17148	47516	Α	17253	1	2393	MEIRGALDLRKRQVLIFLVLLG
						LSRAGTESAHYSVAEETEIGSFV
						ANLARDLGLGVEELSSREARV
						VSDDNKKYLHLDLLTGNLLLN
						EKLDRDELCGSTEPCVLHFQVV
						LENPLOFFRFELCVKDINDHSPT
						FLDKEILIKISEGTTVGATFLME
						SAQDLDVGSNSLQNYTISPNSH
						FYIKIPDSSDRKIYPELVLDRAL
						DYEQEAELRLTLTAVDGGSPPK
						SGTTLVLIKVLDINDNAPEFPQS
						LYEVQVPEDRPLGSWIATISAK
						DLDAGNYGKISYTFFHASEDIR
						KTFEINPISGEVNLRSPLDFEVIQ
	!					SYTINIQATDGGGLSGKCTLLV
						KVMDINDNPPEVTISSITKRIPE
						NASETLVALFSILDQDSGDNGR
						MICSIQDNLPFFLKPTFKNFFTL
						VSEKALDRESQAEYNITITVTDL
						GTPRLKTEYNITVLLSDVNDNA
						PTFTQTSYTLFVRENNSPALHIG
						SVSATDRDSGTNAQVNYSLLPP
						QDRHLPLASLVSINADNGHLFA
						LRSLDYEALQEFEFRVGATDRG
						SPALSSEALVRVLVLDANDNL\
						PFVLYPLQNGSAPCTELVPRAA
						EPGYLVTKVVAVDGDSGQNA
						WLSYQLLKATEPGLFGVWAHN
						GEVRTARLLSERDAAKHRLVV
						LVKDNGEPPRSATATLHVLLVD
						GFSQPYLPLPEAAPAQAQADLL
						TVYLVVALASVSSLFLLSVLLF
17149	47517	A	17254	89	1283	
17150	47518	Α	17255	1	276	
17151	47519	Α	17256	2	259	WQGGILGSDPTPPLTSPNLLQT
						ACFREERDV/RRERGQPLGDHS
						ALCLPRRGVPVPCDGLLCWWG
1	1000	<u> </u>		(0.1	10.5.5	PPDAAEPLRGPSPARAGPVLPG
17152	47520	A	17257	624	4055	
17153	47521	A	17258	3	1784	
17154	47522	A	17259	1	1272	
17155	47523	A	17260	3	576	
17156	47524	A	17261	1	1017	
17157	47525	A	17262	1	1107	
17158	47526	A	17263	256	634	
17159	47527	В	17264	76	1340	133

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
17160 17161	47528 47529	A	17265	1659	540	NLLCCTCSPFSSPVSAPQTCTSN CIALLVHQQQWLVSSHLQKHF LHLHIFLA\LIVFNFSLHSF*IFDF DPWKLPP*PISPTLSSLPSNRHIA LILPCSCMTEDFDFKMPPAVAR YMTSPQAVHIIAVCPFKASRKV SHSSLKIWLFKSV KFFSLRMLNINPHCLLACRVSA ERSAVSLMGFPLCISRPFSLAAL NIFSFISTLVNLMIISLGVALLEE
						YLCGILCIS*IGMLACLARLGKF SWIIR
17162	47530	Α	17267	4880	5121	
17163	47531	A	17268	468	602	CLQLCSFGLGLTW*CGLFFYST *TLK*FFPIL*RMMVVF*WGLH
17164	47532	Α	17269	359	462	
17165	47533	A	17270	457	600	LWWFNALFLCWLTSFLTTTLLE LHECLPVWRD*TRRHLTGWSK GSCTN
17166	47534	A	17271	2405	2830	TPVFRFIALSYTFADSDGEYF*IL LSENPQRFFKCCFNMRSNKIPW DTFANSIKGAEFSGKHFITEFISS SESESESESTRFRDFRGLETSSSL SLDSFSLLGDDFLSFLLSFTIDA VSSKRLSKEIRKARLCWRFPRI
17167	47535	Α	17272	1	2172	
17168	47536	В	17273	446	625	
17169	47537	В	17274	1	2604	
17170	47538	Α	17275	1	3039	
17171	47539	В	17276	1	1383	
17172	47540	Α	17277	1	993	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
17173	47541	A	17278		2494	RSTRQKVNKDTQELNSALHQA DLIDIYRTLHPKSTEYTFFSAPH HTYSKIDHIVGSKTLLSKCKRTE IITNYLSDHSAIKLELRIKNLTQS RSTTWKLNNLLLNDYWVHNE MKAEIKMFFETNENKDTTYQN LWDAFKAVCRGKFIALNAYKR KQERSKIDTLTSQLKELEKQEQ THSKASRRQEITKIRAELKEIET QKTLQKINESRSWFFERINKIDR PLARLIKNKREKNQIDTIKNDK GDITTNPTEIQTTIREYYKHLYA NKLENLEEMDTFLDTYTLPRLN QEEVESLNRPITGSEIVAIINSLP TKKSPGPDGFTAEFYQRYKEEL VPFLLKLFQSIEKEGILPNSFYE ASIILIPKPGRDTTKKENFRPISL MNIDAKILNKILANRIQQHIKKL IHHDQVGFIPGMQGWFNIHKSI NVIQHINRAKDKNHMIISIDEEK AFDKIQQPFMLKTLNKL\G/DKI PRNPTYKGCEGPLQGELQTTAQ GNKRGHKQMEEHSMLMGRKN QYHENGHTAQG/IL*IQCHPHQ ATNDFLHRIGKNYFKVHMEPK KSPHRQVNPKPKEQSWRHHTT *LQTILQGYSNQNSMVLVPKQR YRSMEQNRALRNNAAYLQLSD L*QP*EKQAMGKGFPI**MVLG KLASHM*KAETGSLPYTLYKN
						QFKMD*RFKR*T*NHKNPRRKP RHYHSGHRHGQGLHVQNTKSN
17174	47542	Α	17279	285	502	
17175	47543	Α	17280	1	3189	
17176	47544	В	17281	1	1235	
17177	47545	Α	17282	1	3139	
17178	47546	В	17283	1	1779	
17179	47547	В	17284	1	3127	
17180	47548	A	17285	1	3325	
17181	47549	A	17286	1	3145	
17182	47550	A	17287	1	1095	MIMGDFNTPLSTLDRSTRQKVN KDIQELNSALHQADLIDIYRTLH PKSTEYTFFSASHLTYSKIDHIV GSKALLSNCKRTEIITNYLSDHS AIKLELKIKKLTPNRSTTWKLN NLLLNDYWVHNEMKAEIKMFF ETNENKDTTYQNLWDTFKAEC RGKFIALNAHKRKQERSKIDTL TSQLKELEKQEQTHSKASRRQE ITKIRAELKEMETQK\TLQK\INE SRG/WKKFLEKLCFIELS*ARRI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
17183	47551	A	17288	11	3229	
17184	47552	A	17289	11	1205	
17185	47553	A	17290	1	986	
17186	47554	Α	17291	1	3170	MGDFNTPLSTLDRSMRQKVKK
						DTQELNSALHQADLIDIYRTLH
						PKSTEYTFFSAPHHTYSKIDHIL
						GSKALLSKCKRTEIITNYLSDGS
:					·	AIKLELRIKNLTQNRSTTWKLN
E		1				NLLLNDYWVPNEMKAEIKMFF
						ET\NENKDTTYQNLWDTFKAV
						CRGKFIALNAHKRKQERPKIDT
				1		LTSQLKELEKQEQTHSKASRRQ
						EITKIRAELKEIETQKTLQKIN*S
ŀ		1		-		QSWFFERVKKIDRPLARLIKKK
		1				REKNQIDTIKND
17187	47555	В	17292	1	2148	
17188	47556	Α	17293	1	1416	
17189	47557	Α	17294	1	3099	MGELITPLSTLDRSTRQKVNKD
		1				TQELNSALHQGDLIDIYRTLHP
				1		KSTEYTFFSAPHHTYSKIDHILG
				}		SKALLSKCKRTEIITNYLSDHSA
		1				IKLELRIKNLTQNRSTTWKLNN
						LLLNDYWIHNEMKAEIKMFFET
		ŀ				NENKOTTYQNLWDAFKAVCR
İ				1		GKFIALNAHKRKQERSKIDTLT
						SQLKELEKQEQTHSKASRRQEI
1						TKIRAELKEIETQKTLQKINESR SWFFERINKIDRPLARLIKKKRE
						KNQIDTIKNDK
17190	47558	A	17295	1	3345	KNQIDTIKNDK
17191	47559	В	17296	1	3122	
17191	47560	A	17297	1	2118	
17193	47561	A	17298	965	4091	
17194	47562	В	17299	15	2762	
17195	47563	A	17300	1	3049	MTVKVRITVTSGKKPPWLIPRQ
						TGYGVDLWQTPTDLQLRVLTV
						RRKTNKQKGHPHQNPICTSPSS
						KTKDRSTRQKVNKDIQELNSAL
						HQEDLIDIYRTLHPKSTEYTFFS
						APHHTYSKIDHIVGSKALLSKW
						KRTEITANCLSDHSAIKLELRIK
		1				KLTQNWSATWKLNNLLLNDC
		1				WVHKEMKAEIKMFFETK\ENK\
						DTTYQNLWDAFKA\VCRGKFIA
						LNAHKRKQERSKIDTLTSQLKE
						LEKQEQTHSKASRR

SEQ ID	SEQ ID NO:		SEQ ID NO:			Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	deletion, \=possible nucleotide insertion)
17196	47564	I A	17301	3	4101	
						ALHQVDLIDIYRTLHPKSTEYTF
				1		FSVPHHNYSKIDHIVGSKALLS
						KCKRTEIITNCLSDYNAIKLELR
						TEKLTQNRSTTWKLNNPLLND
						YWVHKEMKAEIKMFFETNENK
						DTTYQNLWDAFKAVCRGKFIA
						LNAHKRKQERSKIDTLTSQLKE
						LEKQEQTHSKATRRQEITKTRA
						ELKKIETQKTLQKINESRSWFFE
						KTNKIDRLLATLIKKKREKNQI
						GARKNDEGAITTN
17197	47565	Α	17302	1	3234	MANKNNQCREVLKGPDELKT
						MAREPHDKCTSFSSQFDQLEER
						VSAMEDQMNEMKREEKFREK
						RIKRNEQRLQEIWDYVKRPNLC
						LIGVPESDGENGTKLENTLQDII
						QENFPNLARQANIQIQEIQRMP
		İ				QRYSLRRTTPRHIIVRFTKVEM
		1				KEKMLRAAREKGRVTHKGKPI
		1				RLTADLSAETLQARRESTRQKV
		1		1		NKDIQELNSAQHQADLTDNYR
			†			TLHPKSTEYTLFSAPHHTYSKID
17100	177566	 	17202	1	3390	HIVGSKALLSKCKRTEI
17198	47566	A	17303		3390	MLTLNHNRLTDAILNISTCLSR HLDLNTGGKKRISYKRNYLME
		l				EHLEGLCSNEFTWENSTIMTDS
						QMLMHREEPGSLPNIPILSIVFT
						NNRSKRQKVNKDIQELNSALH
	}					QADLIDIYRTLHPKSTEYTFFSA
						PHHTYSKTDRIVGSKALLSKCK
						RTEIITNCLSDHSAIKLELRIKKL
						TONCSTTWKLNNLLLNDYWV
						HNEMKAEIKMFFETNENKDTT
						YQNLWDTFKAVCRGKFIALNA
						HKRKQERSKIDTLTS
17199	47567	В	17304	1	2427	
17200	47568	Α	17305	1	4257	
17201	47569	Α	17306	1	3726	MVVKTNARELRDECTSLSSPFN
Ì		ļ				QLEERVSVMENQMNKMKQEE
						KFREKRIKRNEQSLQEIWDYVK
		1.				RPNLCLIGVTESDGENGTKLEN
						TLQDITQENFPNLARKANIQIQE
						TQRTPQRYSSRRATLRHIIVRFT
1						KVEMKEKMLRAAREKDFTPTK
						IKRDKEGHYIMVKGSIQQEELTI
		1				LNIYAPNTGAPRFIKQVLSDLQ
						RDLDSHTLIMGDFNTSLLTLDR
						SMRQKVNKDTQELNSSLHQAD
17000	17570	 	17207	-	1050	LIDIYRTLHPKSIE
17202	47570	Α	17307]]	1059	<u> </u>

	of peptide sequence	hod	in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
17203	47571	Α	17308	130	1092	
17204	47572	Α	17309	1	924	
17205	47573	Α	17310	1	1206	
17206	47574	В	17311	1	1257	
17207	47575	Α	17312		685	MEIILRICKLTISKELLDSGSSRL TMTHYKHRYFSHSAEEQIISEEL NESEMICQHHHTAENDFDELRE EGFRRSVITNFSKLKEDVQTHF KEAKNLEKRLDEWLTRINSVEK TLNDLM*LKTVA*ELRDTYTSF NSRFDQVEERVSVIEDQMNEM EREEKFREKRV*RNEQS/LQEIW NYVKRPNLHLIGVSEIDRENGT KLENTLQDIFQENFTYLARQAN IQIQEI
17208	47576	В	17313	70	1227	
17209	47577	В	17314	640	751	
17210	47578	A	17315		2328	MGVSGAQPGRSCWHRSSTGSQ GLGPVPEPADCGGEAAGRARE EGTVSLTMGTADSDEMAPEAP QHTHIDVHIHQESALAKLLLTC CSALRPRATQARGSSRLLVASW VMQIVLGILSAVLGGFFYIRDY TLLVTSGAAIWTGAVAVLAGA AAFIYEKRGGTYWALLRTLLTL AAFSTAIAALKLWNEDFRYGYS YYNSACRISSSSDWNTPAPTQSP EEVRRLHLCTSFMDMLKVGGQ GEGAAAGLVQNPSGHALGCLD SAASGISDPSVAVLLENVPNQR GESLRCVPVYLGQWNCSKETG MDSGQFGEEKGAHDPSPTYYH ACLSTQWWPSPESPYPQPSGPQ VKEPRNPGKSISMLCLWIRYHA ADSGSSRAKLPLEERLRPFSPTH DCLAFPLPLEKRPEGNVGSRCP SGTKLPEGSGSNICCSAIFPVL QPPLVIPRQTRSGVDLQQTPTD LQLRDLTVRRKTNKQKGHPHQ NPMCTSPSSKTKELKEDVRTHH KEAKNLEKRLDKM\VNRINSVE KTLNDPMELKT\QARELHDKCT SFSSRFDQLEERVSVTEDQMNE MKREEKFREKRVK\RNEQSLQE IRDYVKRPNLCLTGVPESDGEN ETKLENTLQDNIQENFPNLARQ ANIQIQEIQRTPQRYSSRRATPR HIIVRFTKVEMKEKMLRAAGE KGQVTHKGKPITLTADLLAETL
17211	47579	A	17316	508	666	
17211	47580	A	17310	622	780	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
		ŀ		sequence		
17213	47581	В	17318	1	777	
17214	47582	A	17319		508	MSGFEKQYKKIRLDKTLDSENE GLELPVPAKRSNLGVRLYACC GLLLCPAYPQHFAHGYVDKIPG YPGRAGTLTGLHPMQVCRCRR QAPCLKSNNALIVILGTVTLDA VGIGLVMPVLPGLLRDIVHSDSI ASHYGVLLALYALMQF\LCAPK YKLPSENTTNTSMQIN
17215	47583	В	17320	310	1047	
17216	47584	Α	17321	1	1101	
17217	47585	Α	17322	577	735	
17218	47586	Α	17323	337	642	
17219	47587	A	17324	2	653	
17220	47588	A	17325	2	935	
17221	47589	A	17326	532	729	
17222	47590	A	17327	1	675	
17223	47591	A	17328	1	684	
17224	47592	A	17329	2	641	
17225	47593	A	17330		808	
17226	47594	A	17331	562	675	
17227	47595	A	17332	2	1687	DTGNNEWKQQEGLSHNNWLG
						LGCLFGPETPIFYPATLGHTSSIY YERKKEDKNGMKEEHDTSVA MFFTLFNFQISVENIRKGPMVV GESIAQSYLNVVQEIFRSRNHQ LLQDLTRDLCDNLKTCHTSHGS VMAETAVINHKKRKNSPRIVQS NDLTEAAYSLSRDQKRMLYLF VDQIRKSDGTLQEHDGICEIHV AKYAEIFGLTSAEASKDIRQAL KSFAGKEVVFYRPEEDAGDEK GYESFPWFIKQRSNLGVRLYAC CGLLLCPAYPQHFAHGYVDKIP GYPGRAGTLTGLHPMQVCRCR RQAPCMKSNNALIVILGTVTLD AVGIGLVMPVLPGLLRDIVH\SD SIASHYGVLLALYALMQFLCAP VLGALSDRFGRRPVLLASLLGA TIDYAIMATTPVRGHSVLKGKQ HKSVENLQPDNVIEKKIPFSGEK FKPAAEICKSNRELNVNHQDNG ENVSRACQRPLLQPLPSQAWRF RRKNWFCGPGPWSPFSVQSRD LVPCVPAAPALTKRGQGTAQA VASEGGSPKPWQLPHGVEPAG
17228	47596	В	17333	12	60	AQKSRAEVWEPPPRFQKM
17229	47597	В	17334	253	2103	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
17230	47598	Α	17335	469	2512	PTSAYERGQRFASRLQMNLLEY
						FLRWLMAKIFPQAPCMKSNNA
						LIVILGTVTLD\AVGIGLVMPVL
						PGLLRDIVHSDSIASHYGVLLAL
	1					YALMQFLCAPVLGALSDRFGR
	1					RPVLLASLLGATIDYAIMATTP
						VLWIYPLVNVSKSDDTLKINGV
						EDHKTIFDGDGKTYQNVQQFID
						EGNYTSGDNHTLRDPHYVEDK
						GHKYLVFEANTGTENGYQGEE
		ĺ				SLFNKAYYGGGTNFFRKESQKL
	-					QQSAKKRDAELANGALGIIELN
	}			•		NDYTLKKVMKPLITSNTVTDEI
		ĺ				ERANVFKMNGKWYLFTDSRGS
		l				KMTIDGIKLNDIYMLGFVSNSL
						TGPYKPLNKTGLVLQMVLDPN
						DVTFTNFYLPGPQAKANMWLL
1						QAHDKKSFSRIKGTFGPAL
17231	47599	Α	17336	2393	2945	
17232	47600	Α	17337	980	1732	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
17222	47(01	<u> </u>	17338	1	2400	L GRPFPRPPRLCORRLRGGGRMK
17233	47601	A	1/338	L	2400	_
		1				MTGDFEECLKDSPRFRAALEEV
						EGDVAELELKLDKLVKLGIAMI
						DTGKAFCVANKQFMNGIRDLA
						QYSSNDAVVETSLTKFSDSLQE
						MINFHTIL/L*PNSEINLRHSFSN
						FVKEDLRKFKDAKKQFEKVSE
						EKENALVKNAQVQRNKQHEVE
		İ				EATNILTATRKCFRHIALDYVL
						QINVLQSKRRSEILKSMLSFMY
						AHLAFFHQGYDLFSELGPYMK
		ŀ				DLGAQLDRLVGDAAKEKREME
						QKHSTIQQKDFSRDDSKLKYNV
		ł				DAANGIVMEGYLFKRASNAFK
						TWNRRWFSIQNNQVVYQKKFK
						DNPTVVVEDLRLCTVKHCEDIE
						RRFCFEVVSPTKSCMLQADSEK
						LRQAWIKAVQTSIATAYREKG
						DESEKLDKKSSPSTGSLDSGNE
						SKEKLLKGESALQRVQCIPGNA
						SCCDCGLADPRWASINLGITLCI
						ECSGIHRSLGVHFSKVRSLTLDT
						WEPELLKLMCELGNDVINRVY
						EANVEKMGIKKPQPGQRQEKE
		l				AYIRAKYVERKFVDKYSISLSPP
						EQQKKFVSKSSEEKRLSISKFGP
						GDQVRASAQSSVRSNDSGIQQS
						SDDGRESLPSTVSANSLYEPEG
						ERQDSSMFLDSKHLNPGLQLYR
						ASYEKNLPKMAEALAHGADVN
			Í			WANSEENKATPLIQAVLGGSLV
						TCEFLLQNGANVNQRDVQGRG
17234	47602	A	17339	1	782	Teel EEQHOMITTIQUE V QUITO
17234	47603	A	17340	1	475	
17236	47604	A	17341	1	852	
17237	47605	A	17342	432	696	
17238	47606	A	17343	3	296	ETEIEALKEELLFLKKNQEEEV
1,7230	17000					KGLQAQIAISGLTVEVDVPKSQ
		ļ				DLAKIMADIWAQ*DELEDGEDF
						SLRDALDSSNSMQTIRKTTRPI
[VSGWQSGV
17239	47607	A	17344	2	292	SQKKGLVGALTLNPPHFCPCHL
11239	7,007	\frac{1}{2}	11344	[1272	*EVKGLQAQIASS\GLTVEVDAP
						KSQDLAKIMADIRAQYDELAR
						,
						KNREELDKYWSQQVREGKGM
17240	47609	1	17345	1	908	GCQGVGGRQTE
17240	47608	Α	1/343	11	1200	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	1	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
17241	47609	A	17346		1452	TGPEFHDRLRVRRRPQSLSPVL LLSSPDSMSFTTRSTFSTNYRSL GSVQAPSYGAGPRRAARPASY AGAGGSGSRISVSRSTSFRGGM GSGSL\ATG\IAGG\LAGMG\GIQ NEKETMQSLNDRLASYLDRVR SLET\ENRRL\ESKIRE\HLEKKG PQ\VRDWSHYFKI\IEDLRA\QIF\ ANTV\DNARIVL\QI\DNAPSCVR DDFR\VKY\ETELA\MRQSCWRT DIHGAPQGPFDDTNYHTDLQL/ EKTEIRGSSRRELLFQ*RRNHEE /EKLKGLQAQIAQFLGLTRGRL DCPQIFRDLRQDSWADIRGPIID ELASEEPRGSLDKY\WSQQI\EE STTVVTTQF\AEVEAAETT\LTE LRRTVQSLEI\DLDSMRNLKAS\ LENSLREG*RPAYALQ\MEQLN GI\LLHLESELAQTRAEGQRQA QEYEALLNIKVKLEAEIATYRR LLEDGEDFNLGDALDSSNSMQ TIQKTTTRRIVDGKVVSETNDT
17242	47610	A	17347	467	843	
17243	47611	Α	17348	114	1363	
17244	47612	Α	17349	2	281	
17245	47613	Α	17350	1	588	
17246	47614	Α	17351	184	354	
17247	47615	С	17352	391	657	
17248	47616	Α	17353	1	2010	
17249	47617	A	17354	2	443	VPDLPPSHRGFRAAQGGLPPKN KVAALPSWGSANQNSSARWEL VSPPIHRGGS*GPGETGDLDGSS IQRAVELRLVQAFVVPKSSLRD PSREGRGGWAG*VQTKGA*AR PTLLWAPKR/DRGGKAAPTSTW NSPKAGCPSARSSPAPP
17250	47618	Α	17355	l	1359	
17251	47619	Α	17356	260	380	
17252	47620	Α	17357	153	273	
17253	47621	A	17358	270	484	AKYRGTGSSRKALGAHWVPKQ PIPAWQHKDPSGGRPEEQGVKL YREKEFSS*TL*/RI*TGPDAWA YAWADAW
17254	47622	Α	17359	2	941	
17255	47623	Α	17360	1	615	
17256	47624	Α	17361	911	1472	
17257	47625	Α	17362	2	442	
17258	47626	В	17363	76	235	
17259	47627	A	17364	38	190	
17260	47628	Α	17365	134	523	
17261	47629	В	17366	1	1776	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
17262	47630	Α	17367	394	554	
17263	47631	Α	17368	22	831	
17264	47632	С	17369	1	1191	
17265	47633	Α	17370	199	647	PSVPTEGSRNRTTAVAVEERQS DGSGSLSPGKLQTTTTGII\PALL HITGAPVSSDFRLDWLIERPKG GRRAELSIRSLALPLPQHCGSD NDWIPLQLRLLSRASFHSDDDL GFGAYRRLALHILCIHSSKEALE LNTFQPVTFLNAFSK
17266	47634	Α	17371	1	2697	
17267	47635	В	17372	25	1698	
17268	47636	A	17373	2	218	ASLRSSPFSRSLRCQGKNTPERS GLPKALFCTKPSR\ADLGPAGH RRPLPSVQPG*AIPGRVSRGEKN LTNR
17269	47637	Α	17374	2	2543	
17270	47638	A	17375	198	403	PQALREHSPSV*TGRQDRQLLF KDCTCQRQRRDVDSVQGKRRL HISSPQGLNSFPAVSLKLEHNV KAE
17271	47639	Α	17376	58	543	VLYRSSRMCRKIRVLISRPFSIV KICERVLSQTSQGRHDYAAAA VSKTGQKRIHSSGTRGSGAPHS PGWWEPSPDRVGVSICFLFSLK YRLFSSCLVQPCLGPVLGNNSC SPEGPREPSPPSHGLGFPKMCT* TQVPASTTQSDNGQNHPNPGCP MPISPDS
17272	47640	A	17377	14	701	
17273	47641	A	17378	91	309	GRKQFPHLATGKEGCVGGQRL PCVTVELIFPAPAGAAVPRRP*L PRTAPS*LHCLPVCRMPGEAPT ARHAAWV
17274	47642	Α	17379	2	2625	
17275	47643	Α	17380	1	1752	
17276	47644	Α	17381	3257	3412	
17277	47645	A	17382	1	1091	
17278	47646	Α	17383	233	1148	
17279	47647	Α	17384	1	1716	
17280	47648	С	17385	162	356	

17281 47649 A 17386 836 2106 AKKFRKWPLEEQEQ	SEQ ID S	EQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
17281 47649 A 17386 836 2106 AKKFRKRWPLEEQEQ AAPVTALRKRVTSQRK DAVGCAPAQRYTPPS, LGLPLT/TCGARGSWE PLPCRTKHTSGSPRIK FIKNMRQYDTRNSRIV SLCAAFSVLPYGEGUR DSQKQRHPSGGVSVSS EGYELGADGKVSVSVA NALVTHKSDSHGIT CGFHKRVLIFASSHGIT CGFHKRVT CGFHKRVLIFASSHGIT CGFHKRVLI			hod	ŀ	i .	codon for last amino acid	*=Stop codon, /=possible nucleotide
17281 47649 A 17386 836 2106 AKKFRKWPLEEQEQ	St	equence		09/540,21/		or peptide sequence	deterion, \=possible nucleotide insertion)
AAPVTALRKRVTSQRC DAVGCAPAQRYTPPS/ LGLPLT/TCGARGSWE LGLPLT/TCGARGSWE LGLPLT/TCGARGSWE PLPCRTKHTSGSPRHK FIRNMRQYDTRNSRIV SLCAAFSVLPYGEGLR DSQKQRHPSGGVSVS EGVELGADGKVSVA NALVTHKSDSHGLLT RTLPGSRHKPAPTKSA GSDVGDTLEYNPNLLI CGKHRRVLIFASYMTT PSDLKKDMNETTREK! LSKIRSLKREMRSLSE TVAMAYVYFEKLVLQ NRKLCAGACVLLAAK SGVTQLIDKLEERRFF! FEFTVLVALELALYLP 17282 47650 A 17387 1 2531					sequence		
DAVGCAPAQRYTPPS/ LGLPLITCGARGSWE PLPCRTKHTSGSPHK FIKNMRQYDTRNSRIV SLCAAFSVLPYGEGER DSQKQRHPSGGVSVSS EGVELGADGKVVSYA NALVTHKSDSHGLLP RTLPGSRHKPAPTKSA GSDVGDTLEYNPNLLI CGKHKRVLIFASYMTI PSDLKKDMNETFRSIL LSKIRSLKREMRSLSEE TVAMAYVYFEKLVLQ NRKLCAGACVLLAAK SGVTQLIDKLEFRRFI FEFTVLVALELALYLP 17283 47651 A 17388 1 1506 17284 47652 A 17389 34 576 17285 47653 A 17390 112 1840 EELRVREHVTGGICGG DLLGATTLELVAVAP AGGKNLKSPHKVEVD LRWNRSDESVGNVTF TGMDNWIKLSGCQNI SSLKLNVYEEIKLRIR SWYEVJDSFTPFRAQ LEAEDKAIVHISPGTK ALDGLSFTYSLLIWKN RIENIYSRHKIYKLSPE VKAALLTSWKIGVYSI TVENELPPPENIEVSVC LKWDYTYANMTFQVC LKRNFGNHLYKWKQI KTTQCVFPQNVFQKGI ASDGNNTSSWSEEIKF LLPPYFNIRSLDSSFHII SGNTPVIQDYPLIYEIIE NAERKIIEKKTDVTVP VYCVKARAHTMDEKI SDAVCEKTRPGNTSKI SDAVCEKTRPGNTSKI CLALFALFFVIYAAKVI VFFPSLKPSSSIDEYFSI	17281 4	17649	A	17386	836	2106	AKKFRKRWPLEEQEQNVPDGG
LGLPLT/TCGARGSWE							AAPVTALRKRVTSQRCSLEFLE
LGLPLT/TCGARGSWE							DAVGCAPAQRYTPPSAPSSTTP
FIKNMRQYDTRNSRIV SLCAAFSVLPYGEGUED DSQKQRHPSGGVSVSS EGVELGADGKVVSYA NALVTHKSDSHGLPT RTLPGSRHKPAPTKSA GSDVGDTLEYNPNLLI CGKHKRVLIFASYMTT PSDLKKDMMETEREKI LSKIRSLKREMRSLSEE TVAMAYVYFEKLVLQ NRKLCAGACVLLAAK SGVTQLIDKLEERFRFF FEFTVLVALELALYLP T7284 47652 A 17389 34 576					•		LGLPLT/TCGARGSWEEKSDLC
SLCAAFSVLPYGEGLR DSQKQRHPSGGVSVSYA NALVTHKSDSHGLLPI RTLPGSRHKPAPTKSA GSDVGDTLEYPIPNLLL CGKHKVLIFASYMTT PSDLKKDMMETFREKI LSKIRSLKREMRSLSEE TVAMAYVYFEKLVLQ NRKLCAGACVLLAAK SGVTQLIDKLEERFRFI FEFTVLVALELALYLP 17282 47650 A 17387 I 2531 17283 47651 A 17388 I 1506 17284 47652 A 17389 34 576 17285 47653 A 17390 112 1840 EELRVREHVTGGICGG DLLGATTLELVAVAPV AGGKNLKSPHKVEVD LRWNRSDESVGNVTFI TGMDNWIKLSGCQNI SSLKLNVYEEIKNYEEIKNY RIENIYSRHKIYKLSPE VKAALLTSWKIGVYSI TVENELPPPENIEVSVQ LKWDYTYANMTFQW LKRPPGNHLYKWKQI KTTQCVFPQNVFQKGI ASDGNNTSFWSEEIKFI SGNTPVIQDYPLIYEIIF NAERKIIEKKTDVTVP VYCVKARAHTMDEKI SDAVCEKTKPONTSKI CIALFALPFVIYAAKVI VFFPSLKPSSSIDEYFSI CIALFALPFVIYAAKVI VFFPSLKPSSSIDEYFSI							PLPCRTKHTSGSPRHKGLKKTH
SLCAAFSVLPYGEGLR DSQKQRHPSGGVSVSYA NALVTHKSDSHGLLPI RTLPGSRHKPAPTKSA GSDVGDTLEYPIPNLLL CGKHKVLIFASYMTT PSDLKKDMMETFREKI LSKIRSLKREMRSLSEE TVAMAYVYFEKLVLQ NRKLCAGACVLLAAK SGVTQLIDKLEERFRFI FEFTVLVALELALYLP 17282 47650 A 17387 I 2531 17283 47651 A 17388 I 1506 17284 47652 A 17389 34 576 17285 47653 A 17390 112 1840 EELRVREHVTGGICGG DLLGATTLELVAVAPV AGGKNLKSPHKVEVD LRWNRSDESVGNVTFI TGMDNWIKLSGCQNI SSLKLNVYEEIKNYEEIKNY RIENIYSRHKIYKLSPE VKAALLTSWKIGVYSI TVENELPPPENIEVSVQ LKWDYTYANMTFQW LKRPPGNHLYKWKQI KTTQCVFPQNVFQKGI ASDGNNTSFWSEEIKFI SGNTPVIQDYPLIYEIIF NAERKIIEKKTDVTVP VYCVKARAHTMDEKI SDAVCEKTKPONTSKI CIALFALPFVIYAAKVI VFFPSLKPSSSIDEYFSI CIALFALPFVIYAAKVI VFFPSLKPSSSIDEYFSI							FIKNMRQYDTRNSRIVLICAKR
EGVELGADGKVVSYA NALVTHKSDSHGLPP RTI-PGSRHKPAPTKSA GSDVGGTLEYNPNLI CGKHKRVLIFASYMTT PSDLKKDMNETFREKI LSKIRSLKREMRSLSE TTVAMAYVYFEKLVLQ NRKLCAGACVLLAAK SGVTQLIDKLEERFRF FEFTVLVALELALYLP 17282 47650 A 17387 I 2531 17283 47651 A 17388 I 1506 17284 47652 A 17389 34 576 17285 47653 A 17390 112 1840 EELRVREHVTGGICGG DLLGATTLELVAVAPV AGGKNLKSPHKVEVD LRWNRSDESVGNYTF: TGMDNWIKLSGCQNI* SSLKLNVYEEIKLRIKA SWYEVJDSFTPFRAQ LEAEDKAIVHISPGTK ALDGLSFTYSLLIWKN RIENIYSRHKIYKLSPE VKAALLTSWKIGVYSI TVENELPPPENNETVSVC LKWDYTYANNTFQVG LKRNPGGHLYKWKGQI KTTQCVFPQNVFQKGI ASDGNNTSFWSEEIKF LLPPVFNIRSLSDSFHIY SONTPVIQDVPLIYEIIE NAERKIIEKKTDVTVP VYCVKARAHTMDEKI SDAVCEKTKPGNTSKI CIALFALPFVIYAAKVI VFFPSLKPSSSIDEYFSI							SLCAAFSVLPYGEGLRISDLRV
EGVELGADGKVVSYA NALVTHKSDSHGLPP RTI-PGSRHKPAPTKSA GSDVGGTLEYNPNLI CGKHKRVLIFASYMTT PSDLKKDMNETFREKI LSKIRSLKREMRSLSE TTVAMAYVYFEKLVLQ NRKLCAGACVLLAAK SGVTQLIDKLEERFRF FEFTVLVALELALYLP 17282 47650 A 17387 I 2531 17283 47651 A 17388 I 1506 17284 47652 A 17389 34 576 17285 47653 A 17390 112 1840 EELRVREHVTGGICGG DLLGATTLELVAVAPV AGGKNLKSPHKVEVD LRWNRSDESVGNYTF: TGMDNWIKLSGCQNI* SSLKLNVYEEIKLRIKA SWYEVJDSFTPFRAQ LEAEDKAIVHISPGTK ALDGLSFTYSLLIWKN RIENIYSRHKIYKLSPE VKAALLTSWKIGVYSI TVENELPPPENNETVSVC LKWDYTYANNTFQVG LKRNPGGHLYKWKGQI KTTQCVFPQNVFQKGI ASDGNNTSFWSEEIKF LLPPVFNIRSLSDSFHIY SONTPVIQDVPLIYEIIE NAERKIIEKKTDVTVP VYCVKARAHTMDEKI SDAVCEKTKPGNTSKI CIALFALPFVIYAAKVI VFFPSLKPSSSIDEYFSI							DSQKQRHPSGGVSVSSEMVFEL
NALVTHKSDSHGLPT							EGVELGADGKVVSYAKFLYPT
GSDVGDTLEYNPNLLI							NALVTHKSDSHGLLPTPRPSVP
CGKHKRVLIFASYMTT PSDLKKDMNETFREKI LSKIRSLKREMRSLSEE TVAMAYVYFEKLVLQ NRKLCAGACVLLAAK SGVTQLIDKLEERFFF FEFTVLVALELALYLP 17283 47651							RTLPGSRHKPAPTKSAPASTEL
PSDLKKDMNETFREKI LSKIRSLKREMRSLSEI TVAMAYVYFEKLVLQ NRKLCAGACVLLAAK SGVTQLIDKLEERFRF FEFTVLVALELALYLP 17282							GSDVGDTLEYNPNLLDDPQWP
LSKIRSLKREMRSLSEE TVAMAYVYFEKLVLQ NRKLCAGACVLLAAK SGVTQLIDKLEERFRF FEFTVLVALELALYLP							CGKHKRVLIFASYMTTVIEYVK
TVAMAYVYFEKLVLQ NRKLCAGACVLLAAK SGVTQLIDKLEERFRIF FEFTVLVALELALYLP 17282							PSDLKKDMNETFREKFPHVKLT
NRKLCAGACVLLAAK SGVTQLIDKLEERFRF FEFTVLVALELALYLP 17282							LSKIRSLKREMRSLSEECSLEPV
NRKLCAGACVLLAAK SGVTQLIDKLEERFRF FEFTVLVALELALYLP 17282	1						TVAMAYVYFEKLVLQGKLIKQ
T282							NRKLCAGACVLLAAKISSDLRK
T282							SGVTQLIDKLEERFRFNRRDLIG
17282							FEFTVLVALELALYLPENQVLP
17284 47652 A 17389 34 576 17285 47653 A 17390 112 1840 EELRVREHVTGGICGG DLLGATTLELVAVAPV AGGKNLKSPHKVEVD LRWNRSDESVGNVTF: TGMDNWIKLSGCQNIT SSLKLNVYEEIKLRIA SWYEV\DSFTPFRKAQ LEAEDKAIVHISPGTK ALDGLSFTYSLLIWKN RIENIYSRHKIYKLSPE VKAALLTSWKIGVYSI TVENELPPPENIEVSVC LKWDYTYANMTFQVG LKWDYTYANMTFQVG KTTQCVFPQNVFQKGI ASDGNNTSFWSEEIKF LLPPVFNIRSLSDSFHIY SGNTPVIQDYPLIYEIIF NAERKIIEKKTDVTVP VYCVKARAHTMDEKI SDAVCEKTKPGNTSKI CIALFALPFVIYAAKVI VFFPSLKPSSSIDEYFSI	17282 4	17650	Α	17387	1	2531	
17285 47653 A 17390 II2 1840 EELRVREHVTGGICGG DLLGATTLELVAVAPV AGGKNLKSPHKVEVD LRWNRSDESVGNVTFSTGMDNWIKLSGCQNITSSLKLNVYEEIKLRIRASWYEVDSFTPFRKAQ LEAEDKAIVIHISPGTK ALDGLSFTYSLLIWKN RIENIYSRHKIYKLSPEVKAALLTSWKIGVYSITVENELPPPENIEVSVOLKWDYTYANMTFQVOLKRNPGNHLYKWKQIKTTQCVFPQNVFQKGIASDGNNTSFWSEEIKFLLPPVFNIRSLSDSFHISGNTPVIQDYPLIYEIIFNAERKIIEKKTDVTVPVYCVKARAHTMDEKISDAVCEKTKPGNTSKICIALFALPFVIYAAKVIVFFPSLKPSSSIDEYFSI	17283 4	17651	A	17388	1	1506	
DLLGATTLELVAVAPV AGGKNLKSPHKVEVD LRWNRSDESVGNVTFS TGMDNWIKLSGCQNIT SSLKLNVYEEIKLRIRA SWYEV\DSFTPFRKAQ LEAEDKAIVIHISPGTK ALDGLSFTYSLLIWKN RIENIYSRHKIYKLSPE VKAALLTSWKIGVYSI TVENELPPPENIEVSVQ LKWDYTYANMTFQVQ LKRNPGNHLYKWKQI KTTQCVFPQNVFQKGI ASDGNNTSFWSEEIKF LLPPVFNIRSLSDSFHIY SGNTPVIQDYPLIYEIIF NAERKIIEKKTDVTVPI VYCVKARAHTMDEKI SDAVCEKTKPGNTSKI CIALFALPFVIYAAKVI VFFPSLKPSSSIDEYFSI	17284 4	17652	A	17389	34	576	
AGGKNLKSPHKVEVD LRWNRSDESVGNVTFS TGMDNWIKLSGCQNIT SSLKLNVYEEIKLRIRA SWYEV\DSFTPFRKAQ LEAEDKAIVIHISPGTK ALDGLSFTYSLLIWKN RIENIYSRHKIYKLSPE VKAALLTSWKIGVYSI TVENELPPPENIEVSVQ LKWDYTYANMTFQVQ LKRNPGNHLYKWKQI KTTQCVFPQNVFQKGI ASDGNNTSFWSEEIKF LLPPVFNIRSLSDSFHIY SGNTPVIQDYPLIYEIIF NAERKIIEKKTDVTVPI VYCVKARAHTMDEKI SDAVCEKTKPGNTSKI CIALFALPFVIYAAKVI VFFPSLKPSSSIDEYFSI	17285 4	17653	A	17390	112	1840	EELRVREHVTGGICGGSQMMV
LRWNRSDESVGNVTF: TGMDNWIKLSGCQNITSSLKLNVYEEIKLRIRA SWYEV\DSFTPFRKAQ LEAEDKAIVIHISPGTK ALDGLSFTYSLLIWKN RIENIYSRHKIYKLSPE VKAALLTSWKIGVYSH TVENELPPPENIEVSVQ LKWDYTYANMTFQVQ LKRNPGNHLYKWKQI KTTQCVFPQNVFQKGI ASDGNNTSFWSEEIKF LLPPVFNIRSLSDSFHIY SGNTPVIQDYPLIYEIIF NAERKIIEKKTDVTVP VYCVKARAHTMDEKI SDAVCEKTKPGNTSKI CIALFALPFVIYAAKVI VFFPSLKPSSSIDEYFSI							DLLGATTLELVAVAPWVMSAA
TGMDNWIKLSGCQNIT SSLKLNVYEEIKLRIRA SWYEV\DSFTPFRKAQ LEAEDKAIVIHISPGTK ALDGLSFTYSLLIWKN RIENIYSRHKIYKLSPE VKAALLTSWKIGVYSI TVENELPPPENIEVSVQ LKWDYTYANMTFQVQ LKRNPGNHLYKWKQI KTTQCVFPQNVFQKGI ASDGNNTSFWSEEIKF LLPPVFNIRSLSDSFHIY SGNTPVIQDYPLIYEIIF NAERKIIEKKTDVTVPI VYCVKARAHTMDEKI SDAVCEKTKPGNTSKI CIALFALPFVIYAAKVI VFFPSLKPSSSIDEYFSI							AGGKNLKSPHKVEVDIIDDNFI
SSLKLNVYEEIKLRIRA SWYEV\DSFTPFRKAQ LEAEDKAIVIHISPGTK ALDGLSFTYSLLIWKN RIENIYSRHKIYKLSPE VKAALLTSWKIGVYSI TVENELPPPENIEVSVQ LKWDYTYANMTFQVQ LKRNPGNHLYKWKQI KTTQCVFPQNVFQKGI ASDGNNTSFWSEEIKF LLPPVFNIRSLSDSFHIY SGNTPVIQDYPLIYEIIF NAERKIIEKKTDVTVPI VYCVKARAHTMDEKI SDAVCEKTKPGNTSKI CIALFALPFVIYAAKVI VFFPSLKPSSSIDEYFSI							LRWNRSDESVGNVTFSFDYQK
SWYEV\DSFTPFRKAQ LEAEDKAIVIHISPGTK ALDGLSFTYSLLIWKN RIENIYSRHKIYKLSPE VKAALLTSWKIGVYSI TVENELPPPENIEVSVQ LKWDYTYANMTFQVQ LKRNPGNHLYKWKQI KTTQCVFPQNVFQKGI ASDGNNTSFWSEEIKF LLPPVFNIRSLSDSFHIY SGNTPVIQDYPLIYEIIF NAERKIIEKKTDVTVPI VYCVKARAHTMDEKI SDAVCEKTKPGNTSKI CIALFALPFVIYAAKVI VFFPSLKPSSSIDEYFSI							TGMDNWIKLSGCQNITSTKCNF
LEAEDKAIVIHISPGTK ALDGLSFTYSLLIWKN RIENIYSRHKIYKLSPE VKAALLTSWKIGVYSI TVENELPPPENIEVSVO LKWDYTYANMTFQVO LKRNPGNHLYKWKQI KTTQCVFPQNVFQKGI ASDGNNTSFWSEEIKF LLPPVFNIRSLSDSFHIV SGNTPVIQDYPLIYEIIF NAERKIIEKKTDVTVPI VYCVKARAHTMDEKI SDAVCEKTKPGNTSKI CIALFALPFVIYAAKVI						ľ	SSLKLNVYEEIKLRIRAEKENTS
ALDGLSFTYSLLIWKN RIENIYSRHKIYKLSPE VKAALLTSWKIGVYSI TVENELPPPENIEVSVO LKWDYTYANMTFQVO LKRNPGNHLYKWKQI KTTQCVFPQNVFQKGI ASDGNNTSFWSEEIKF LLPPVFNIRSLSDSFHIV SGNTPVIQDYPLIYEIIF NAERKIIEKKTDVTVP VYCVKARAHTMDEKI SDAVCEKTKPGNTSKI CIALFALPFVIYAAKVI VFFPSLKPSSSIDEYFSI							SWYEV\DSFTPFRKAQIGPPEVH
RIENIYSRHKIYKLSPE VKAALLTSWKIGVYSH TVENELPPPENIEVSVO LKWDYTYANMTFQVO LKRNPGNHLYKWKQI KTTQCVFPQNVFQKGI ASDGNNTSFWSEEIKF LLPPVFNIRSLSDSFHIV SGNTPVIQDYPLIYEIIF NAERKIIEKKTDVTVP VYCVKARAHTMDEKI SDAVCEKTKPGNTSKI CIALFALPFVIYAAKVE							LEAEDKAIVIHISPGTKDSVMW
VKAALLTSWKIGVYSI TVENELPPPENIEVSVO LKWDYTYANMTFQVO LKRNPGNHLYKWKQI KTTQCVFPQNVFQKGI ASDGNNTSFWSEEIKF LLPPVFNIRSLSDSFHIY SGNTPVIQDYPLIYEIIF NAERKIIEKKTDVTVPI VYCVKARAHTMDEKI SDAVCEKTKPGNTSKI CIALFALPFVIYAAKVI							ALDGLSFTYSLLIWKNSSGVEE
TVENELPPPENIEVSVO LKWDYTYANMTFQVO LKRNPGNHLYKWKQI KTTQCVFPQNVFQKGI ASDGNNTSFWSEEIKF LLPPVFNIRSLSDSFHIY SGNTPVIQDYPLIYEIIF NAERKIIEKKTDVTVPI VYCVKARAHTMDEKI SDAVCEKTKPGNTSKI CIALFALPFVIYAAKVE							RIENIYSRHKIYKLSPETTYCLK
LKWDYTYANMTFQVO LKRNPGNHLYKWKQI KTTQCVFPQNVFQKGI ASDGNNTSFWSEEIKF LLPPVFNIRSLSDSFHIY SGNTPVIQDYPLIYEIIF NAERKIIEKKTDVTVPI VYCVKARAHTMDEKI SDAVCEKTKPGNTSKI CIALFALPFVIYAAKVE							VKAALLTSWKIGVYSPVHCIKT
LKRNPGNHLYKWKQI KTTQCVFPQNVFQKGI ASDGNNTSFWSEEIKF LLPPVFNIRSLSDSFHIY SGNTPVIQDYPLIYEIIF NAERKIIEKKTDVTVPI VYCVKARAHTMDEKI SDAVCEKTKPGNTSKI CIALFALPFVIYAAKVE							TVENELPPPENIEVSVQNQNYV
KTTQCVFPQNVFQKGI ASDGNNTSFWSEEIKF LLPPVFNIRSLSDSFHIY SGNTPVIQDYPLIYEIIF NAERKIIEKKTDVTVP VYCVKARAHTMDEKI SDAVCEKTKPGNTSKI CIALFALPFVIYAAKVF		{					LKWDYTYANMTFQVQWLHAF
ASDGNNTSFWSEEIKF LLPPVFNIRSLSDSFHIY SGNTPVIQDYPLIYEIIF NAERKIIEKKTDVTVP VYCVKARAHTMDEKI SDAVCEKTKPGNTSKI CIALFALPFVIYAAKVF							LKRNPGNHLYKWKQIPDCENV
LLPPVFNIRSLSDSFHIY SGNTPVIQDYPLIYEIIF NAERKIIEKKTDVTVPI VYCVKARAHTMDEKI SDAVCEKTKPGNTSKI CIALFALPFVIYAAKVE VFFPSLKPSSSIDEYFSI							KTTQCVFPQNVFQKGIYLLRVQ
SGNTPVIQDYPLIYEIIF NAERKIIEKKTDVTVPI VYCVKARAHTMDEKI SDAVCEKTKPGNTSKI CIALFALPFVIYAAKVI VFFPSLKPSSSIDEYFSI			1	ŀ			ASDGNNTSFWSEEIKFDTEIQAF
NAERKIIEKKTDVTVPI VYCVKARAHTMDEKI SDAVCEKTKPGNTSKI CIALFALPFVIYAAKVI VFFPSLKPSSSIDEYFSI							LLPPVFNIRSLSDSFHIYIGAPKQ
NAERKIIEKKTDVTVPI VYCVKARAHTMDEKI SDAVCEKTKPGNTSKI CIALFALPFVIYAAKVI VFFPSLKPSSSIDEYFSI							SGNTPVIQDYPLIYEIIFWENTS
SDAVCEKTKPGNTSKI CIALFALPFVIYAAKVI VFFPSLKPSSSIDEYFSI							NAERKIIEKKTDVTVPNLKPLT
CIALFALPFVIYAAKVI VFFPSLKPSSSIDEYFSI							VYCVKARAHTMDEKLNKSSVF
VFFPSLKPSSSIDEYFSI							SDAVCEKTKPGNTSKIWLIVGI
VFFPSLKPSSSIDEYFSI							CIALFALPFVIYAAKVFLRCINY
							VFFPSLKPSSSIDEYFSEQPLKNL
, , i i i i i i i i i i i i i i i i i i			1				LLSTSEEQIEKCFIIENISTIATVE
1 1 1 1 1 1						}	ETNQTDEDHKKYSSQTSQDSG
1 1 1 1 1 1							NYSNEDESESKTSEELQQDFV
17286 47654 A 17391 3 373	17286 4	47654	A	17391	3	373	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	,
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
17287	47655	A	17392	1	432	LPRFVGQQPRPLGPLRGSGSEG
						CRAPGEAVRAPDYAFPIPGTRA
İ						S/PLNRFPLGGTAWPQAHPTNP
		ŀ				KPAPAPGAQLDDPPRYSRHGLS
						RWGLPLQSLVLLPPLQRPMLPQ
						AAAPNTPWWPWTLRAGRREN
						AHAQRKLHLWAGKVE
17288	47656	Α	17393	98	376	
17289	47657	Α	17394	1	364	
17290	47658	Α	17395	1	1230	
17291	47659	A	17396	892	2004	RRMSFMSSEVLVGELMSPFDQS
		1				GLGA\EESLGLLDDYL\EVAKHF
						KPHGFSSDKAKAGLLPNGLAV\
		ł				DGLVSPSNNSKE\DAF\SGTDW
						DVGRKWVLKEFDFGCPCLGIE*
						PWETIARLTFLTTFGIDICDSLLP
		ŀ				P*FQGRLIKAAPPRRLNPIG\HLP
						QKFLTKPRPGLAPFTF\LQPL\PL
		l				SPGVL\SSTPDSFPFSFRSWGS*K
	Ì					WDIH*KGD*/RKPDYTAYVA\MI
						PQCIKE\EDTPSD\ND\SGI\CMSP
						ES\YLGSPQHSPST\RGSPNRSLP
						SPGVLCGSARPKPYDPPGEKMV
						AAKVKGEKLDKKLKKMEQNK
						TAATRYRQKKRAEQEALTGEC
						KELEKKNEALKERADSLAKEIQ
	15.00	ļ	15005		1200	YLKDLIEEVRKARGKKRVP
17292	47660	Α	17397	43	1300	DSCCIQGTMRLSRTQQRPGVEG
				Ì		SLKQHQRLHSNFSSINESVFIVS
				ŀ		FSLNTIGSGGRGAEGEP/GREPL
]				SRPQPPAGPLGGALPAPP\SLMP
1		ļ				QL*SE*GRLLQLLYLLLQ/PAVC
					}	SCSRCSAPRSRCVARPAARTGL PTPAPASSP/CSRSLSSSRSLSSSC
						RVSCCHSLSSCCRSLSCSRSSST
					ł	SSRSSSTSSSSLSLSSSLSCSSCRL
						SCSNSFSTSSSSLSLSSSLSCSSC CLSCSNSFSTSSSSLSLSSSLSCS
						SCRLSCSNSFSTSSSSLSCSGSFS
					:	CSHSLSSCSRSLSSCSHSLSSCSR
						SLSSCSHSLSSCSRSLSSCSHSLS
İ						SCSRSLSSCSHSLSSCSRSLSSCS
						CSLSSCSRSLSSCSHSLSSCSHSL
						SSCSRSLSSCSCSLSSCSRCLSSC
						SCSLSSCSRSLSSCLLFRRLNI
17293	47661	A	17398	288	574	SCOLSGCSROLSGCLLI RRCINI
1/293	14/001	<u> </u>	1/390	1200	214	

SEQ ID	SEQ ID NO:	i	SEQ ID NO:	<u> </u>	L .	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
17294	47662	A	17399	230	648	VIAPGLWMMAADIPRVTTPLSS LVQVPQEEDRQEEEVTTMILED DSWVQEAVLQEDGPESEPFPQS AGKGGPQEEVTRGPQGALGRL RELCRRWLRPEVHTKEQMLTM LPKEIQAWLQEHRPESSEEAAA LVEDLTQTL*DSWVQEAVLQE DGPESEPFPQSAGKGGPQEEVT RGPQGALGRLRELCRRWLRPE VHTKEQMLTMLPKEIQAWLQE HRPESSEEAAALVEDLTQTL
17295	47663	Α	17400	105	381	SSGFWPLRFPLDVIKAVSTIAVH EKEESLWPRVAVFSTLAP/ESPP WGEAPQSAGR*SGVPEDHVHL RYRGTLSLACCSSRHHPPAQLS HYKN
17296	47664	A	17401	1023	1915	
17297	47665	A	17402	3	277	QVLLQGALLLLITVAQAGRVLK LGVSREFQDLLILWGKETFRR/C PGPCSPALHLQPGSSIP/SSQQLL *YLAAEQPQLHRGPPRTPAPVS GAE
17298	47666	Α	17403	3	299	
17299	47667	С	17404	46	398	
17300	47668	A	17405	171	403	GKNSTLQQFNNLLHYSSITVAV NNPTLGNSVSVFPLYILFTGIHQ LVKVILRKLGK*RFGLELASLSL LFLPPVSLF
17301	47669	Α	17406	291	467	
17302	47670	A	17407	2	395	DHAYVFLGRTACYLMKAYRK VNPRVIFSSNTIETHPKDLLHSC DHPFAEKTQFTVSTLDDVKNSG SIRDNYCRTSEISAVHIDTECVS VMLQAGTPPLQVK*RKNFQQR KQG*KMKWDPSDISNSMAEVL
17303	47671	A	17408	1	149	EKEIPGPWLLGAMRFRFCGDLD CPDWVLAEISTLAK\IPL*SCGCS AARY
17304	47672	A	17409	555	1156	QFYNKTPSRSECPNLLHHMRTE GPPGGVDQPDGHQAGWPPPEP AGQQRGRQSGCHLLQGGASPR PSCGTVHARACTPPASCPPCHR YRSHLQRGKWWPK/GLPQHTR LDHTEVLGAAEGSTVSAVPSAT RAARDHVRRLQTRKCFCKGAL LLLITAARAGRVLFGQPQLLQF TGQGFAIDTVLGRTGEDETQHC HCGLHIA
17305	47673	A	17410	3	366	
17306	47674	A	17411	408	1702	
17307	47675	Α	17412	1	2661	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
17308	47676	A	17413	2	2288	
17309	47677	Α	17414	1	1992	MGDFNTPLSTLDRSMRQKVNK
						DIQELNSAVHQVDLIDIYRTLHP
	1					KSTEYTFFLAPHRTYSKIDHIVG
1	-	l				SKALLSKCKRTEIIRNCLSDHSA
		Ì				IKLELRINKLTQNCSTTWKLNN
		1				MLLNDYWVHNEMKAEIKMFF
					•	ETNENKDTTYKNLWDTFKAVC
		ł		ļ		RGKFIALNAHKRKQERSKIDTL
						TSQFKELEKQEQTHSKASRRQE
						ITKIRAELKEIETQKTLQKINESR
						SWFFEMISKIDRPLARLIKKKRE
						KNQIDTIKNDKGDIITDPTEIQTT
						IREYYKHLYAIKLENLEEMDKF
						PYTNTLPRLNEEEIESLNRTITGS
						EIEAIIKSLPTKKSPGPDGFTAKF
						YQRYKEELVPFLLKLFQSIEKE
						GILPNSFYEASIILIPKP/DVGSSS
		1				QGNQAGERNKGYSFRKRGSQI
		İ				VPVCR*YDCTFRKSHFLSPKSP*
						ADKQLQQSLRIQNQCAKITSIPI
						HHQQTNREPNHE*TPIHNCFKE
						NKIPRNPTYKGCEGPLQGELQT
						TAQ*NKKGHKQMEEHSMLMD
						RKNQYRENGHTAQGNLQIQCH
						PHQATNDFLHRIGKNYFKVHM
						EPKKSPHCQDTPKQKE*SWRH
						HAT*LQTILQGYSNQNSMVLVP
						KQRYRPMEQNRALRNNTTHLQ
						QSDL*QT*QKEEMGKGFPI**M
						VLGKLASHM*KAETGSLLYIL
17310	47678	Α	17415	1	1677	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			•	sequence		
17311	47679	Α	17416	1	2178	MGDFNTPLSTLGRSTRQKVNK
.,						DTQELNSALHQADLIDIYGTLH
						PKSTEYTFFSAPHHTYSKIDHIV
		i				GSKALLSKCKRTEIITNCLSDHR
						AIKLELRIKKLTQNRSTTWKLN
:		1				NLLLNDYWVHNEMKAEIKMFF
		l	,			ETNENKDTTYQNLWDTFKAVC
	-					RGKFIALNAHKRKQERSKIDTL
						TSQLKELEKQEQTHSKASRRQE
						ITKIRAELKEIETQKTLQKINESR
]		l				SWFFERINKIDRPLARLIKEKRE
		l				KNQIDATKNDKGDITTDPTEIQ
						TTIREYYKTLYANKLENLEEMD
						KLLDTYTLPRLNQEEVESLNRPI
		l				TGSEVVAIINSLPTKNSPGPDGF
						TAKYYQRYKEELVPFLLKLFQS
					-	TEKEGILPNSFYEASIILIPKPGR
						DTTKKENFRPISLMNVDAKILN
						KILANRIQQHIKKLIHHDQVGFI
						PGMQGWFNICKSINVIQHINRT
						KDKNHMIISIDAEKAFDKIQQPF
						MLKTLNKLVLEVLARAIRQEKE
						IKGIQLGKEEVKLSLFADDMIV
						YLENPIISAQNLLKLISNFSKVS
						GYKINVQKSQAFLYTNNRQTES
						QIMSELPFTIASKRIKYLGIQLTR
						DVKDLFKENYKPLLNEIKEDTN
						KWKNIPCSWIGRINIVKMAILPK
]		VIYRFNAIPIKLPMTFFTELEKTT
						LKFIWNQKRARITKSILSQKNK
						AGGITLPDLKLYYKAIVT\KTA
						WYCYQNRDIDQWNRTEPSEIM
17312	47680	Α	17417	1	1770	
17313	47681	Α	17418	1	1101	
17314	47682	Α	17419	1	1614	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
17315	47683	Α	17420	l	1878	MVLALTGNKADLASKRPLEFQ
						EAQAYADDDSWLFMETSAKG
						AMNVNEIFMAIAEKLPKNKPQ
						NAAGAPDRNQGYLSACEFSGR
İ						KLPKQTTKHIKQSLRTKLCGAS
						YASSNLIPRKLYEPPLVIPRKTG
						SGVDLQQTPTDLQLWVLTVRR
						KSNKQKGYPHQNPICMLPSSKT
		1	,			KDFKPTKIKRDKDGHYIMVKG
						SIQQEELTVLNINAPNTGAPRFI
						KQALRDLQRDLDSHTIIMGDFN
						TPLSTLDRSTRQKVNKDIQELN
		İ				SALHQADLIDIYRTLHPRSKDY
		1				TFFSAPHHTYSKIDHIVGSKALL
						AKCKRTEITRNCLSDHSAIKLEL
						RIKKLTQKCSTTWKLNNLLLND
		1				YWVHNEMKAEIKMFFETNENQ
						DTTYQNLWDTFKAVCRGKFIA
						LNAHKRKQERSKIDILTSQLKE
		ł				LEKQEQTHSKASRRQEITKIRVE
						LKEIKTQKTLQKINESRSWLFE
		1				KINKIDRQLARLIKKKREKNQID
]				AIKNDKGDITTDPTEIQTTIREY
						YKHLYAKKLENLEEMDKFLDT
						YTLPRLNQEEVESLNRPITGSEI
						EAIIKSLPAKKSPGPDGFTAEFY
					•	QRCEKELAQFLLRLFKSTEKEGI
						LPNSFYEDSIIL/ILPKPGRDTTK
						KENFRPISL
17316	47684	Α	17421	1	1761	
17317	47685	Α	17422	1	1875	
17318	47686	Α	17423	1	1611	
17319	47687	Α	17424	1	4539	
17320	47688	В	17425	1	6294	
17321	47689	A	17426	2	1913	
17322	47690	A	17427	1	1026	
17323	47691	A	17428		2112	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		İ		sequence		
17324	47692	A	17429	1	1884	MENGFDELREEGFRRSNYSELQ
						EEIQTKGKEVKNIEKNLDECITR
		ŀ				IPNTEKCLKKLMELKTKARELR
]	EECISLRSRCDQLEESVSVMED
						EMHEMKREGKFREKRIKRNEQ
						SLQEIWDYVKRPNLRLIGVPED
						RSMRQKVNKDTQELNSALHQA
		ļ				DLIDIYRTLHPKSTEYTFFSAPH
		ŀ]			HTYSKIDHIVGSKALLSKCKRT
		ŀ				EIITNYLSDHSAIKLELKIKKLTQ
		l				NFSTTWKLNNLLLNDYWVHN
]			KMKAEIKMFFETNENKDTTYO
1						i
						NLWDTFKAVCRGKFIALNAYK
						RKQERSKIDTLTSQLKELEKQE
						QTHSKASRRQEITKIRAELKEIE
						TQKTLQKINESRSWFFERINKID
						RPLARLIKKKREKNQIDAIKND
ŀ						KGDITTDPTEIQTTIREYYKHLY
		ĺ				TNKLENLEEMDKFLDTYTLPRL
						NQEEVESLNRPITGSEIVAIIHSL
						PTKKSPGPDGFTAQFYQRYKEE
						L/RNKIPRNPTYKGCEGPLQGEL
						QTTAQGNKRGDKQMEEHSML
						MGRKNQYRENGHTAQGNLQIQ
		ŀ				CHPHQATNDFLHRIGKNYFKV
						HMEPKKSPHRQVNPKPKEQSW
1						RHHAT*FQIILQGYSNQNSMVL
ļ		l				VPKQRYRSMEQNRALRNNATY
		<u>L</u> _				LQLSDL*QT*EKQAI
17325	47693	Α	17430	1	1695	
17326	47694	Α	17431	1	3779	MELKTKARELREECRSLRSRRN
						QLEERVSAMEDEMNEMKREG
						KFREKRIKRNEQSLQEIWDYVK
						RPNLRLIGVPESDAENGTKLEN
						TLQDIIQEDFPNLARQANVQIQE
		ŀ			1	IQRTPQRYSSRRATPRHIIVRFT
						KVEMKQKMLRAAREKDFKPT
						KIKRDKEGHYIMVKGSIQQEEL
					1	TILNIYAPNTGAPRFIKQVLSDL
}					1	QRDLDSHTLIMGDFNTPLSTLD
		1				RSTRQKVNKDTQELNSALHQA
		[DLIDIYRTLHPKSTE
17327	47695	A	17432	1	735	
17328	47696	Α	17433	1	2541	
17329	47697	Α	17434	1	1461	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
17330	47698	A	17435	2	429	AACAAAMSLVIPEKFQHILRVL
						NTNIDGRRKIAFAITASSSSSSS
						SSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSS
		l		1		SSSSSSSSVNLLSNGLYL/VKG
						VGRRYAHVVLRKADIDLTKRA
						GELTEDEVERVITIMQNPRQYKI
						PDWFL
17331	47699	Α	17436	1	615	
17332	47700	В	17437	264	556	
17333	47701	Α	17438	1	406	FFFFFLQKVGIYLHIKTFQQHRL
		ŀ				WVTSAAFLSLSSMAAVGTSSSL
						SENSGPGHGGKQTF*GP/EGTDI
						PEGRHHFSAQAFSVVAGRGTA
		İ				AALLHGSHVPQGR\SRIGSRSGG
						PPSHPDTCSGSSHSAGRSCPHSC
						TG
17334	47702	Α	17439	3	3484	
17335	47703	В	17440	42	367	
17336	47704	В	17441	203	382	
17337	47705	Α	17442	22	266	YRDLPHFLLNSDLIFNMSTSLRË
						TMILIRTLSRVPL/PLHGVIQPIS
						KIQGLVLNALNQVQESIFQVSF
						NLFPDAFQHVIRA
17338	47706	A	17443	1	1137	
17339	47707	Α	17444	1218	1482	QSPWPSLLRGDVSITPNLIFSCL
						SIGWPFCLV/SFC*PKK*TFQGV
						GSEVYDVKHFASCVLCHKQMN
						KLNLYFRNTYFVALLSTELFEN
17340	47708	Α	17445	2	377	
17341	47709	Α	17446	1	211	
17342	47710	Α	17447	2	732	CSSPSN/TILGVQKLNSQWRLA
						QDFRLINEAVIPLYPVVPNPYTL
						LSQIPEEAESFTVLDLKDAFFCIP
						LHSDSQFLFAFEDPTDHNSQLT
						WMVLPQGFRDSPHLFGQAQAQ
						DLVHFSSPGTLVVQYMDDLLL
						ATSSEASCQQATLDLLNFLANQ
						GCKRVGIALGVLTQTHGTTPQP
						VAYLSKETDVAAKGPLTAILLL
						LAFGPCIFNLLVKLVSSRIDAIK
						LQMVLQMEPQMSSTNNFYRGP

SEQ ID			SEQ ID NO:	1	1	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		057570,217	sequence	or peptide sequence	governous, v. possible indeceding insertion,
17343	47711	A	17448	1	3003	MPGAQDLQPTMPEPPHPQWAP
						TWPEPPRRVPPPAPRRPVPLTA
						QGLSVDQSRLGKFSDDPDRYID
						VLQGLGQTFDLTWRDVMLLLD
						QTLAFNEKNAALAAAREFKDT
						WYLSQVNDRMTAEERDKFPSG
						QQAVPSMDPHWDLDSDYGDW
			ļ			SHKYLLTCVLEGLRRIRKKPMN
				:		YSMMSTITQGKEENPAFLEQL
				į.		WEALRKYIPLSPKSLEGQLILKD
						KFITQSAADIRGKLQKQSLVPE
		ļ	15110		1004	QNLEALLNLATSVFYNRDQ
17344	47712	Α	17449	5	624	IQPILAYPHPKTLKQLWGFLGIT
						GFCPVWIPRCSKIAKSLYTLIKE
						TQRANTHLVEWEPEAETAFKT LKQALVQAPTLSLPTGENFSLY
		1				VTERAGIALGVLTQTRGTTPQP
						VTYLKDGEPMKCDCQQIIVQTY
						ATQDDLLEVPLANPDLNLYTD
				1		GSSFVENGIRRAGYAIVSDVTIL
						ESKPLPPGTSAQLAELVALTRA
		1				LELGKGKE
17345	47713	A	17450	636	2628	SNDRTEDDCGKHPFMSSPP\TEP
						WVCLIEGQEIDFLLDTGTTFSV
						LIPCLGRLSSRSVTIQGILGQPVT
						RYFSHLLSCNWETLLFSHAFLV
						MPESPTPLLGRDILAKAGAIISM
						KTGNKLPICCPLLEGINPEVWA
						LEGQFGRAKNAHPLQIRLKDPI
ļ						SFPYQRQYPLRPEAHKGLQDIV
		1				KHLKAQDSVRKCSSPCNTPILG
						VQKLNSQWRLVQDLRLINEAVI
						PLYPVVRNPYTLLSQVPEEAEW
					T .0	FTVLDLKD
17346	47714	Α	17451	2	568	CSSPSN/TILGVQKLNSQWRLA
		l				QDFRLINEAVIPLYPVVPNPYTL
		l				LSQIPEEAESFTVLDLKDAFFCIP
		1				LHSDSQFLFAFEDPTDHNSQLT
						WMVLPQGFRDSPHLFGQAQAQ DLVHFSSPGTLVVQYMDDLLL
	1	1				ATSSEASCQQATLDLLNFLANQ
		1				GCKRVGIALGVLTQTHGTTPQP
1		1				
	<u></u>	1		<u> </u>	<u></u>	VAYLSKETDVAAK

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	Sequence		03/3/3/3/21/	sequence	or popular sequen-	
17347	47715	Α	17452	254	849	LQPESLEIAGILVQ*MIE*QPKK
						GTNSIPVSKPSP/GVQKPNGQW
						RQVQDLRLISDAVIPLYPAVSNP
						YTLLSQILEEAEWFTVLDLKDA
						FFCIPLRSDSQFLLAFEDPTDHT
						SQLTWTVFTQGFMDTPHLFGQ
						SLAQDLGHFSSPGTLVLQYVDD
						LLLAKQQATLDLLNFLANQGY
						KLSKLKAQLCELLVFSSCARMH
17348	47716	Α	17453	1	369	
17349	47717	A	17454	1	1632	
17350	47718	Α	17455	636	1837	SNDRTEDDCGKHPFMSSPP\TEP
						WVCLIIEGQEIDFLLDTGTTFSV
		ł				LIPCLGRLSSRSVTIQGILGQPVT
		l				RYFSHLLSCNWETLLFSHAFLV
						MPESPTPLLGRDILAKAGAIISM
						KTGNKLPICCPLLEGINPEVWA
						LEGQFGRAKNAHPLQIRLKDPI SFPYQRQYPLRPEAHKGLQDIV
	j					KHLKAQDSVRKCSSPCNTPILG
						VQKLNSQWRLVQDLRLINEAVI
						PLYPVVRNPYTLLSQVPEEAEW
						FTVLDLKDAFFCIPLHSDSQFLF
						AFEDPTDHTSQLMWMVLPQRF
						RDSPHLFGQAQAQDLGHFSSPG
		l				TLVLQYSEIAKTLYTLIKEMER
		1				ANTHLVEWEPEAETAFETLKQ
						ALVQAPALSLPTGQNFALYVIE
		1				RAGIALGVLTQTHRTTPQPVAY
17351	47719	A	17456	2	52	EISFVENGIQRAGYAIVSDVTVL
	ĺ					ESKPLLPGTSAQLAELVALTRA
		1				FELGKGKRINANTDSKYAYLIL
	ì	l				HALAAIWKEREFLISGGTPIKYH
		1				KEIMELLHAMQKPKEVAVLHC
	į					QSHQKG*EWDTKGRLCHS
17352	47720	Α	17457	1625	2505	PGVQKP/GQWRLVQDLRLINEA
						VIPLYPVVPKPYTLFSQIPKEAE
						SFTVLDLKDAFFCIPLHSDSQFL
		ĺ				FAFEDPTDHTSQLTWTVLPQRF
						RDSPHLFGQALVQDLGHFSSPG
		1				TLVLQFVDDLLLAASSEASCQQ
		1				ATRDLLNFLANQGYKASKSKA
						QLCLQQVKYLGLILARGTRALS
						KKRIQPILAYPRPKTLKQLRGFL
						GITSFCRLQIPGYSKMARPLYTL
		l				IKETQRANTHLVEWEPEAETAF
						KSLKQALVQAPALSLPTRQNFS
				}		LYVRERAGIALEVLTQTHGTTL
		ļ	1-1	<u> </u>		QPVAKG
17353	47721	Α_	17458	<u> </u>	2415	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
17354	47722	A	17459	2646	5167	KNSGHILGTCPGYRYAKGKET
	:					GKEIIKGPQNPPGYRLCPLQAV
1				1		GGGEFGPTQVHVPFSLSDLKQI
				1		RVDLGKFSDDPDRQYPLRPEAH
						EGLQDILRYLKVQGLVRKCSSP
						CNTRILGVQKPKGQWRLVQDL
						RFINEAVIPLHSVVPNSYTLLSQI
						PEEVEWFTVLDLKDAFFCIPLH
į.						SVCQFLFAFEDPTDHTSQFTWT
						VLPQGFRDSPHLFGQILAQDLG
	ļ					HFSSPGTLVLQYVDDLLLATSL
						EASCQQATLDLLNLLANQGYK
						ASKPKAQLCLQQVKYLDLSLA
		1		ł		RVTRALSKERIQPILACPRPKTL
						KQLRGFLGITNFCQLWIPGYSE
İ						MARPLYTLIKETQRANSHLVER
						EREAETAFKTLKQALVQALALS
						LPTGQNFSLYARERAGIALGILT
1						QTHGTTPQPVTYLRKLIYSFVE
						NGMRRAGYAIVSDVTVPESKPL
						PPGTSTQLAELVVLTQALELRK
						KKRINVNTDSKYAYLILHAHAA
		l				IWKEREFLTSGGTPNKYHKEIM
	1	1				ELLHAVQKPKEVAVIPCQSHQK
		l				GEEK/EAEGNRRADAE\AKIAA
1						RRNLPTEIPMEGPLVWNNPVQE
						IKPQYSLTKTEWGLSQGHSFLP
		i i				LGWLTTLEGKAGRATPNPRNV
						YFPLHLRGLRTGRLPLSQSTRA
		1				QEMTLVRPRPSSSPPPEKAAEG
		1				RGRRCSSETKLRHRRLFRPLRTP
		l				LCATAPLAASVAAVLAAAPAA
17355	47723	В	17460	-1	2208	
17356	47724	Α	17461	2	168	
17357	47725	Α	17462	890	1034	
17358	47726	Α	17463	386	619	TFITGDCLGTKN/HIFDGCGVLS
						EVGLVLDCSNQRFKAGPA*LCQ
			}			SPFKNIL*ESLPMCMCRQLEKH
						AHPQGLCHTQVT

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
17359	47727	A	17464	123	2169	DLGHRGSREMASVA\LEDVAV NFTREEWALLGPCQKNLYKDV
						MQETIRNLDCVGMKWKDQNIE
]	DQYRYPRKNLRCRMLERFVES
				1		KDGTQCGETSSQIQDSIVTKNT
						LPGS/VGPYESRMSGEVIMGHSS
				ļ		LNCYIRVGAGHKPHEYHECGE
						KPDTHKQRGKAFSYHNSFQTH
						ERLHTGKKPYDCKECGKSFSSL
:						GNLQRHMAVQRGDGPYKCKL
						CGKAFFWPSLLHMHERTHTGE
						KPYECKQCSKAFSFYSSYLRHE
						RTHTGEKPYECKQCSKAFPFYS
						SYLRHERTHTGEKPYKCKQCS
						KAFPDSSSCLIHERTHTGEKPYT
		ł				CKQCGKAFSVSGSLQRHETTHS
						AEKPYACQQCGKAFHHLGSFQ
						RHMIRHTGNGPHKCKICGKGF
						DCPSSLQSHERTHTGEKPYECK
						QCGKALSHRSSFRSHMIMHTG
						DGPHKCKVCGKAFVYPSVFQR
						HERTHTAEKPYKCKQCGKAYR
						ISSSLRRHETTHTGEKPYKCTLG
						KAFIDFCSFQNHKTTHTGEKPY
]						ECKECGKAFSRFRYLSRHKRTH
				-		TGEKPYECKTCRKAFGHYDNL
	İ					KVHERIHSGEKPYECKECGKAF
						SWLTCFLRHERIHMREKSYECP
						QCGKAFTHSRFLQGHEKTHTG
]	ENPYECKECGKAFASLSSLHRH
						KKTHWKKTHTGENPYECKECG KAFASLSSLHRHKKTHWKKTH
17260	47720	<u> </u>	17465	1	1217	KAFASLSSLHKHKKIHWKKIH
17360 17361	47728 47729	A	17466	1	1317 336	
17362	47730	C	17467	51	353	
17363	47731	A	17468	190	348	YAIHIMDVHTVRSLYYFNCFRY
17303	17731	``	17 100		3 10	*ACYCCHYLTDIPEITKLSGCSY
						GGTSLIWG
17364	47732	A	17469	1	1716	
17365	47733	Α	17470	1	840	
17366	47734	A	17471	699	1283	LLTARLTWVSARLMSYVLLMC
						*LFPRLLSCLTTPPHCSFSICFVI
						C/FKNVDLKGFEPDVRILLTKYS
						NSNGSQSPWMEEQIRDAGERV
						HSAFSLNWGKPLVFEKDRPYSV
						SVFIGLFLPMAEKTGPLKGNPG
						GEFLLFLRQIQLVWLPAAGIMT
						EHYFPITATQALEINNAISTIMEI
						FPTVSFASQWITASFSV
17367	47735	Α	17472	300	461	
17368	47736	Α	17473	1	2451	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
17369	47737	Α	17474	2	3542	
17370	47738	A	17475	1	1156	
17371	47739	Α	17476	1	1176	
17372	47740	Α	17477	1	257	
17373	47741	Α	17478	295	1353	
17374	47742	Α	17479	3	1772	
17375	47743	Α	17480	2	284	
17376	47744	Α	17481	1	141	
17377	47745	Α	17482	1	506	PHSPLWKTLCSPSSGVCCGFVT
		İ				LFCNDRFACLFQLTPPPSPAWM
						TAAASLLASLLPALPPLNLQDA
1						KFVEERRKQLQNYLRTVMNKV
						IQMVPEFAASPKKETLIQLMPFF
						VMGLQVGMYGLFHFGVMMDQ
						PQPSLTRADESPCDRVLSPCAA
						HRKQRDCCLSWKAEPE*FLVK*
						KTLCSPSSGVCCGFVTLFCNDR
					•	FACLFQLTPPPSPAWMTAAASL
į						LASLLPALPPLNLQDAKFVEER
	•					RKQLQNYLRTVMNKVIQMVPE
						FAASPKKETLIQLMPFFVMGLQ
						VGMYGLFHFGVMMDQPQPSLT
						RADESPCDRVLSPCAAHRKQR
						DCCLSWKAEPE
17378	47746	Α	17483	1	741	
17379	47747	Α	17484	1	993	
17380	47748	В	17485	277	416	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
17381	47749	1	17486	686	2400	TGMSVDEKPDSP\MYVYESTVH
1/381	4//49	Α	17400	000	2400	CTNILLGLNDQRKKDILCDVTLI
1				1		· · · · · · · · · · · · · · · · · · ·
		:				VERKEFRAHRAVLAACSEYFW
						QALVGQTKNDLVVSLPEEVTA
						RGFGPLLQFAYTAK\LLLSRENI
						REVIRCAEFLRMHNLEDSCFSF
						LQTQLLNSEDGLFVCRKDAAC
					İ	QRPHEDCENSAGEEEDEEETM
	ļ					DSETAKMACPRDQMLPEPISFE
						AAAIPVAEKEEALLPEPDVPTD
						TKESSEKDALTQYPRYKKYQL
				1		ACTKNVYNASSHSTSGFASTFR
						EDNSSNSLKPGLARGQIKSEPPS
		<u>.</u>				EENEEESITLCLSGDEPDAKDR
						AGDVEMDRKQPSPAPTPTAPA
						GLCLERSRS/GNYGQPH/VGQK
			•			EVSNFTMGS/PLRGPGLEALCK
						QEGELDRRSVIFSSSACDQVSTS
						VHSYSGGARFLATEHQEPGLM
						GDGMYNQVRPQIKCEQSYGTN
				:		SSDESGSFSEADSESCPVQDRG
						QETGSTSDLRLSTKLQSCSGPM
						FKGPTATVAATESWGTVLQNIP
						LGCAGTGEEVTYTNPPPRGRAA
						LVPRSPLTVYSSSLPEHQTSLKQ
						CAQAGPGVCLLVHNHWQLLEG
17382	47750	A	17487	1	1089	
17383	47751	С	17488	295	348	
17384	47752	Α	17489	1	711	
17385	47753	Α	17490	1	1248	
17386	47754	A	17491	135	259	
17387	47755	Α	17492	434	506	
17388	47756	A	17493	3	227	
17389	47757	C	17494	3	236	
17390	47758	C	17495	25	153	
17370	11770	<u></u>	11770	1~~	1.00	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO:	of peptide sequence	1100	09/540,217	codon for peptide	of peptide sequence	deletion, =possible nucleotide insertion)
	-			sequence		·
17201	1	I _A	117496	 	1557	MASPEAVASQDNDGFSQKPPPT
17391	47759	A	,,,,,,	,	(1337)	PLFASRSVNRLKSQRTPRGEVE
						SVTDGEKTDGSWRMTVDYCKL
			}	}		NQVVTPIAAAVPDVVSLLEQIN
						TSPDITLVHYIDDIMLNGSSEQE
1						VTNTLDLLVRHLCATGWEINPT
1						KIQGSSTSVKFLGVQWCGDCR
1				[DVPSKVKDKLLHLAPPTTKKEA
						QHLTGLFGFRRKYIPYLGVLLC
		}				PIYQVTRKAASFQWRPEQEKAL
	}					QQVQAAMQAALPLGPYDPAGP
						MVLEIAVADTEAVWGH/WHSA
				1		SFPSHPCHCPMGP*TKWP*WOG
						WRLCMGSATWTSTHEG*PGYG
						HQ*VPNLPAAETNTEPAI*HHS/
						SRRSASYLVAG*LYWTSCIMER
						AE\HLLWIWVCLSCMQCFCQD
						YHPWTHGMPYPPLWYSTQLCL
}						*PRHSLYH*RSVAVGSCSWNSL
			}		· · · · · · · · · · · · · · · · · · ·	VLPCSPSS*SSRVDRTVERPFEIT
				1		TTMPTSQESRVQESRGGSGSGT
					1 1	THHHP**STSNIFASCFYGITFC
}	1)	WSRGLSSRGRNAATRRQNNSIK
		1		<u> </u>	<u> </u>	L*VKIATWTLWAPPTSKL
17392	47760	Α	17497	2	1120	PSKVKEKLLHLAPPTIKKEAQR
	[1			LVGQLGFWRQHIPHLGALLWLI
						YQATQKAASFEWGPEQEKALQ QVQVAVQAALPLGPYDPADPM
						VLEVSVADRDAVWSLWQAPIG
			1			ESQQRPLGFWSKALPSSTDNYS/
						CFRCAIIY*FMGCCQ*FGWIVRD
1	1	1				LEEA*LENW*QRNLGKRYVDG
	Ĭ					PL*VVKNWKDICIPCECSPTGDL
			1		j j	SRGGF**SSG*DDPFCGHHSASF
		1			,	PSHPCCRPMGP*TKWPWWQE*
					1 1	RLCMDSATWTSTHQS*PDYSH
1	1				i i	C*VPNLPAAETNTETSIWHHFS
1		1			1 1	G*SASYLLAG*LH*TSSIMEKAE
}	1					VSPHWNKHSLWIWVCLSSMQC
	}					FCQDYHPWTHSMPYPLSWYST
						QHCLSLQRERHSLYR*RSAAVG
						ACS
17393	47761	Α	17498	1	1908	
17394	47762	A	17499	1	915	
17395	47763	A	17500	2	670	_
17396	47764	A	17501	1	1332	
17397	47765	A	17503	151	674	
17398	47766	A	17504	134	825	
17399	47767	A	17505	134	447 778	
17400	47768	Α	17506	98	//0	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
17401	47769	A	17507	1	830	MHCFFIFSAAWFCLWSISLHLV
1,401	1,705	``	1,20,	1	030	ERPDMAVLLFPFGLRLGLMLQ
						CPRGYWPVLLGAEWLLIYWLT
						QAVGLTHFPLLMIGSLLTLLPV
		1				1 7
1						ALISRYRHQRDWRTLLLQGAA LTAAALLQSLPWLRHGKESWN
						1
		1				ALLLTLTG/EPDAGPDMSGVLA
						LSRQ*HLAAARSVTGFSANQLA
						RATSGLVLAAVCYQSLAPVGIA
						GRTVALYAILSGAADYRAGLA
						LWLARGADCDVDERHRADRH
						QSWRDLPWTYCSGDCDSRHGA
<u></u>						QGQKARAPPKHDQFPHAIFRG
17402	47770	Α	17508	2252	2772	RSKTTAFTPLRVRTSAHARPAG
						PAPIIATFLPVCFTPDISGRQPISN
						ALSLI*RSMLPMVTAPNSSFRV
		1				QEPS/SQTVLRADSSANFRQGV
		Ì				GLVRQFRRFKNTPLVGELQPVR
						DVVVYRAFPLAVWVAARQAA
		ļ				VSLRFGLAFGKRLVNFNKLNFA
	<u> </u>					DLQRFLWRINALQVDKLITF
17403	47771	Α	17509	539	1077	
17404	47772	C	17510	192	396	
17405	47773	С	17511	10	294	
17406	47774	A	17512	2	379	KRPGSGITEATSSRRQRPTRRLQ
		ļ				GQLPGQEKRVLKATEGIANPS
		ŀ				WGWGGAVSPLRASLTSLRHKI
	ŀ		<u> </u>			NNHYNTENWKLPIKDS\HLIPSP
						QAPSMPFPLANPPW\APHPREK
						MWNPQIQGTNCIWVCGP
17407	47775	Α	17513	291	590	
17408	47776	Α	17514	187	589	VLSFLIHRWEGDKDSLLGPNFS
	1	1				QTPQPSRRFICAFPWENPS*EPS
			İ			YPSISDPGFLILLPSKGDVWIHL
		1				GPIFSKKPLYS*SSFLVIFYPVTP
						SFLFLGYKFPLAHAVFRVEPTLS
			_	L		LLIQNSHCHGCYTYPDGPE
17409	47777	Α	17515	1509	1748	IGVVREGIPVLCLFSKGMLPVF
						AHSV*YWLWVCHR*LLLF*DTS
						HQYLIY*EFLA*SVVDFCQRPFL
		1			1	HLVLDFFWLVSY
17410	47778	Α	17516	152	568	CQLELSCMKCPSTPVGRSLPVG
						CTRSGTQLRWHSVPWQSWCTV
1		1				LGESLLSGLVVLFRASRPE**NP
1						LKLLPQLPLPPDALSQGDGSFV
						CKHPTGAVTFPSGMPLGSMTIF
						KILILPIHEHGMFFSLSVSSFVSL
			L	L	L	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
	·			sequence		
17411	47779	Α	17517	152	568	CQLELSCMKCPSTPVGRSLPVG
						CTRSGTQLRWHSVPWQSWCTV
		1				LGESLLSGLVVLFRASRPE**NP
		l				LKLLPQLPLPPDALSQGDGSFV
						CKHPTGAVTFPSGMPLGSMTIF
İ		ļ				KILILPIHEHGMFFSLSVSSFVSL
17412	47780	Α	17518	152	479	CQLELSCMKCPSTPVGRSLPVG
						CTRSGTQLRWHSVPWQSWCTV
						LGESLLSGLVVLFRASRPE**NP
						LKLLPQLPLPPDALSQGDGSFV
						CKHPTGAVTFPSGCPTHTLQA
17413	47781	Α	17519	1	474	
17414	47782	Α	17520	1326	1712	FLYVIKALGKKQLQFGQQILIPS
						IELLDYLINPGSGCVSLWKLNK
						PQYCYLIHFSGLTELSWKVHTIL
						LEGHYEFLEYAADR*PICLFSW
						KERQLVSFLDYSLHHLLGSFYV
						VFFFFLYLLVRLCWDL
17415	47783	Α	17521	18	236	
17416	47784	Α	17522	281	1005	TGWSPLQLSKATVARLPLWIPP
						LWAGHLGELWLASGEYPSGLK
		1				LPEEGTGSNLRCSAASAGDTQA
						NRVWSGPAANFSRPAAEGPAF
						HALRGPSVHCSPLCVHEFSSFSS
		ļ	<u> </u>			LISENMQYLVFCSGVRFLRIMVI
			}			SSIHILCLVPKRTRVLGRLSLTC
						FRFLRGPKEPRITIPHSSAPPLGH
						VIAIPLP*SFMKPSPEADAPMTG
						SFLKSPSEADTGATLPVQSVVPI
17417	17705		17523	1	1524	PLSILPILEVGGVLFT
17417	47785	A		001	1524	SRKMGAPNA**SAWSWSREGL
17418	47786	Α	17524	881	1108	AAP*SMFSSLSADSSADCKLPE
						GKVQPASSTGLTARASKDLYFC
						CCFSISARAS
17419	47787	C	17525	1	867	CCI SISAINAS
17420	47788	A	17526	203	514	SSSTARPSQTSSAEESPTSWYRG
17,120	17700	1	17320	203		DNPSKPMDPLVAGVFGAIVGA
l						ASVLGNAPLDGIETGCGAWSST
						NAEHTGLWLQILRKEGLKDL*N
						GTVPRLGRWTASPWM
		<u> </u>	l	<u>t</u>	<u> </u>	GI TI LEGICW I TO I WITE

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO:	Nucleotide location of first		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO:	sequence	nou	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	Sequence		03/210,211	sequence		,, · · p 0000000 n == 00000000 n == 000000000 n
17421	47789	I A	17527		1578	I MAKKEQLLSAAPSKTNAEGQEI
	.,,,,,,,	-				PSGVYTTRALGFKHKTGQPFGQ
						TPSYPQEFILYTNDTWNASKTE
						PFTPLERGLKPGSQESSGWHLV
						GTPLGRSFQRKEQAAIFAVLQP
						PLVIPRQTGSGVDLQQTPTDLQ
						KRVLTVRRKTNSININKKDDHA
						KTTSKGHQQQRPKVDKSTKMR
						KNQCKKATNSKNQNVSSPPKD
						HNSSTAKEONWMENEFDKLTE
						VGFRSSQFDQVEERISDIEDKLN
					1	EIKHEDKIREKRMKKKEQSLQE
						IWEYVKRPNLRLIGVPESDGEN
						GTKLENTLQDTIQENFPNLARQ
						ANIQIQEIQRTPQRYSSRRATPR
ĺ						HII/E*IHQG*NEGKTVKGSQRE
						K*GQEIPSGVYTTRALGFKHKT
						GQPFGQTPSYPQEFILYTNDTW
ŀ						NASKTEPFTPLERGLKPGSQESS
						GWHLVGTPLGRSFQRKEQAAIF
						AVLQPPLVIPRQTGSGVDLQQT
						PTDLQKRVLTVRRKTNSININK
						KDDHAKTTSKGHQQQRPKVDK
						STKMRKNQCKKATNSKNQNVS
						SPPKDHNSSTAKEQNWMENEF
						DKLTEVGFRSSQFDQVEERISDI
						EDKLNEIKHEDKIREKRMKKKE
			ļ			QSLQEIWEYVKRPNLRLIGVPE
						SDGENGTKLENTLQDTIQENFP
			<u> </u>			NLARQANIQIQEIQRTPQRYSSR
						RATPRHIIGRFTKVEMKEKRLR
		<u> </u>				AVREKSRVTHKGKLIRLTADLS
17422	47790	Α	17528	738	1323	ATGGHSNQRQRS*KLSKKFRR
						MYN*NNQYREVLKGADGAEN
						QGSRTTETLCM*RMRQSLQF\D
:			ļ			SDLVRHQRKHSGEKPYECCEC
İ						GKAFRGSSELIRHRRIHTGEKPY
						ECGEYKSYECIACGKAFGRSST
						LIENQRIHTGEKPYECNECGKSF
						NQSSALTQHQRIHTGEKPYECS
17400	45501	ļ.	17500		656	ECRKTFRHRSGLMQHQRTHTR
17423	47791	Α	17529	1	656	MAGKPEYDKTISTSIVLNALNA
						LGVSAEASGRNDLVVKTVEGD
						RKVSGSAYRETKDRGFHHGTL
						LLNADLSRLANYLNPDKKKLA
						AKGITSVRSRVTNLTELLPGITH
					•	EQVCEAITEAFFAHYGERVEAE
						IISPNKTPDLPNFAETFARQSSW
						EWNFGQAPAFSHLLDERFTWG
						GVELHFDV/GKRPYHPRPGVYR
	<u> </u>	<u> </u>			<u> </u>	QPQPRAAGSPRRTTARLPFLSL

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
17424	47792	A	17530	1046	1401	LLDIRMLCHGKGVIIHVLRLQF LLGRPAPHRLHAALQLPGLSLH GTLALGLWLQLRQLLLLGQPF GISGCGSGCDPEQDCV/DCGRG CWPRKCPQGSHSAPRSRGAPHR PRPQQRAAGRQ
17425	47793	A	17531	112	1170	YPFFHLIDSTIDTCVCFTKFLCC VFQLHQVIYVLL*TGYSLIFDAD DLWMGFLRGHPFC*CYCFLFVS FPSNSQAPLLQVCWSLLEVHSR PCSPGYHQRRLQNSKDCCLLLP LEALSQSGTHQMPAGALLYEM SVNPCWEEHQTDQLFTHESTFI RSKTQKKVPDKLLDSSTVTPLF KITENIGCVMTGMTADSRSQVQ RARYEAANWKYKYGYEIPVD MLCKRIADISQVYTRNAEMRPL GCCMILIGIDEEQGPQVCKCDP AGYYCGCKATAVGVKQTESTS FLEKKVKKKFDWTFEQTLETAI TCLSTVLSIDFKPSEIEVGVVTV ENPKFRILTEAEIDAHLVALAER
17426	47794	A	17532	2244	2432	DGQQLIALHRLALRELQQAVH AGIAPAGENPV*WWFRNRQNP LIALRSLPAFQSGNLSCQLH
17427	47795	A	17533	251	546	SSISDILSSA*SIRLLILVYAS*SS RAVFFSTIRSFIFFSKLVILVSPSC NLLSRFLASFHWVRTCSFSSKEF VIITHLLPTSVNSSYSSLFSFVHF G
17428	47796	A	17534	597	1092	FTNQKESRTRWIHSRILPEVQG GTGTIPSETIPINRKRGNPP*LIL* GQRHSDTKAG\RDTTKKENFRP ISLMNIDAKILNKILANRIQQHI KKLIHHDQVGFIPGMQGWFNIR K/CNKCNPAYKQSQRQK/HT*L SQ*MQKKPLTKFNNPSC*KLSIN *VLVGRISK
17429	47797	A	17535	1	2153	
17430	47798	A	17536	2164	2427	
17431	47799 47800	A	17537 17538	1	627	GPRWGPRVHRAPDPVRIPSSGP LLPPRFYPRYVLP\PPSASTSRPC QRTSASGRCPCPMH/DTVKATM PIWV/DPPVPDHYEGEAEHQGIL VTHPHHQRCPAGHRHRVVF
1/432	7/000	<u> </u>	17330	1	027	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
17433	47801	A	17539	583	1145	GNGRTVDTQVCQAPITSQSLER PADPNP\GEQDAEAGPASRRDT EAERLRFRQFQYHVASGPHLA VGQLWTLCRQWLRPEARSKEQ MLEVLVLEQFLGALPSKMRTW VQSQGPRSCREAASLVEDLTQ MCQQEGLAAHPSHDTAIRGIPR LAASWETASAMEATADARQKR THTEKEAWLLEAPHELA
17434	47802	A	17540	1	1301	
17435	47803	A	17541	1	1232	
17436	47805	A	17542	146	510	MNTANVEKPYVQKYQTAFVLV MKHRQCQLASHLVQNYRS/IQL WAWE\KIQDPGTTCPSFNTITQG SKKPYPDFVARLRDAAQNSITD ENARKVIVELMAYGNANPECQ SATKPLKGKVPTGSDVISDQKA VGVQAAKRRNRTDTGLGRSAR KSGKGVTRMLPKHASMILDGG GPATSPSCCRFL SPYWNKGKSPGWLAES\FKRPV
	47806					FRQGAK\EPYPDFAASLQDAAQ KSITNENARKVIVELMAYENAN PECQSAIKPLKGKVPVGSDVISR GFQGQQPPLSQVLQGMGQLPQ YSNCPRPQAAVQQ
17438		A	17544	2	828	
17439	47807	A	17545	1	420	
17440	47808	A	17546	1	1947	DI KADIMBOOKA EUN THATES
17441	47809	A	17547		399	PLKVNIYPDSKYAFHILHHHER GFLTTQGSSIINASLIKTLLKAA LLPKEARVIHCKGHQRASDPIA QDNAHTDKIAKKAASIPTYTPH FQFFSFSS\ATPTYSPAETSTYQS LPTQGKWFLDKETYFLPASQ
17442	47810	A	17548	1	2382	

SEQ ID	_		SEQ ID NO:	I .		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first	t e	*=Stop codon, /=possible nucleotide
	sequence	ŀ	09/540,21/	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				a cqueine		
17443	47811	Α	17549	1	2194	MAPEQGRVSILQTLFIEYHGYSI
					ļ	FVVKGDLPDCEADQLLQMIRV
						QQMHRPKLIGEELAQLKEQRV
						HKTDLERVLEANDGSGMLDED
						EEDLQRALALSRQEIDMEDEEA
						DLRRAIQLSMQGSSRNISQDMT
						QTSGTNLTSEELRKRREAYFEK
						QQQKQQQQQQQQGDLsGQ
						SSHPCERPATSSGALGSDLALLL
						TQSLFGSLFTWTRVTFGAEDPG
1						QEDSFRRRVPCPCPHSVRRSTY
		ŀ				DLGSSDQPAQGTSHEFQIGLLL
						TRLSQVPILPQPPVPHPIILLSPPP
						HTVSGLQFLSVTSPPLPAQQFLL
	<u> </u>					KEVAGAEGIAKVIPTCPVPLLG
						QDILTKLSASLIIPGLQPHLIATL
					^	LPNPKPPSPLPLVSPNLNPQVW
Ì					i	DTSTPSLATDQTPITIPLKPNHP
						YLAQCHYSIPQLALRGLKPVIT
	ŀ	ŀ				RLLQHGLLKPINSSYNSPILPVQ
	}					KPDKSYRLPQDLDLINQIVLPIH
						PVASPHIIPDTTPDPHDCISLIHL
-	İ					AFTPFPHTSFFPIPHPDHTWFID
-						l i
						GSSTWPNRHSPAKVGYAIVVSS
						TSILEATALPPSTTSQQAELIALT
1					+	RALALAKGLHVNIYTDSKYAF
				1		HILHHNAVIWAERGFLSTQRSSI
-						IIASLIKTLLKAALLPKEAGVIH
		İ				CKGHQKASDPIAQGNAYADKS
						DNGLAFTSQI/I/HKQFLRLLVFS
						GTFIPLTVLNLQERTGQYFYLLP
17444	47010	 	10550	1	255	FTELEPVLEKLQGTVHLNFYMD
17444	47812	A	17550	1	355	
17445	47813	C	17551	99	335	L CDI DEGI CDCOLLICOCATA CO
17446	47814	A	17552	21	447	LGPLPFSLSPCCLHCQGKRLCG
		1				HHEEARRKNVSIPRKEAGIIHC
						KGHQK/A/SDPIAQGNAYADKT
						KFLDLAFPPLRLSFTCQITQAVS
						QALGIQWNLHTLTILNLQKRIR
						ACPQDATGYSPFELLYGCSFLL
	.==.	ļ				GPSLIPDTSPT
17447	47815	A	17553	119	239	TQKGQRAALFLICILSPSPLPTL
		L				GKNFKN*IPALKPHNRI
17448	47816	A	17554	1	798	
17449	47817	В	17555	460	1059	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
17450	47818	Α	17556	3	1476	IPTCPVPLLGQDILTKLSASLIIP
						GLQPHLIATLLPNPKPPSPLPLV
		l		!		SPNLNPQVWDTSTPSLATDQTP
						ITIPLKPNHPYLAQCHYSIPQLA
		l			,	LRGLKPVITRLLQHGLLKPINSS
						YNSPILPVQKPDKSYRLPQDLD
						LINQIVLPIHPVASPHIIPDTTPDP
		l	,			HDCISLIHLAFTPFPHTSFFPIPHP
						DHTWFIDGSSTWPNRHSPAKV
]				GYAIVVSSTSILEATALPPSTTS
		1		}		QQAELIALTRALALAKGLHVNI
		ł				YTDSKYAFHILHHNAVIWAER
						GFLSTQRSSIIIASLIKTLLKAAL
					·	LPKEAGVIHCKGHQK\ASDPIA
						QDNAYADKLAKKAASVPTSVP
						HGISQAPPPLPTHQARYWQIDF
					:	THMPRVRKLKYLLVWVDTFTG
		İ				WVEAFPTGSKKATAVISSLLSDI
						IPQFSLPTSIHSDSRLAFISQITQA
						VSQALGIQWNLHISYRPQSSGK
	•				•	VEQTNGLLKTHLTKLSLQLKK
ŀ						DWTVLLPLALLRIRACP
17451	47819	Α	17557	24	594	VGSGDLPWEGNPLSSCSLLHEK
		l				DPPTTSGPQTDQPKEHLTNFKS
		1				GCSSPGPARSQFFLSLRSSTL*SF
						YHLPSSHLVCLTVSFRD*PSPTC
						PAIDS*KGG/CELKA/RVKEHFC
		ł				HDLPASSMQSNYPVHDSGHLT
						TGPSLFKPTLNIEYSTGHAPYHV
						RSACFPFAFCHDCQFPEASPEAE
						AAAILSVQPAEP
17452	47820	Α	17558	1	1152	
17453	47821	C_	17559	301	462	
17454	47822	Α	17560	1	423	
17455	47823	Α	17561	421	856	KLINLFSNFCSYSRQFLSFSSTL
						YFVWVFCYNRCCFSK*RMIVHI
						TNRDINCQIAYARIEGDMIVCA
						AYAHELPKYGVKVGLTNYAAT
						YCTGLLLAHRLLNRFGMDKIC
						YLDAGLARTTTGNKVFGDLDG
						GLSIPQSTK*FPGYD

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
17456	47824	A	17562	3	1082	GPLCIALALQEFGTRRRWACRS LSSSGRRSLFRRMGSVKAVKNK AYFKRYQVKFRRRREGKT\DYY ARKR\LVIQDKNKYNTPKY\RMI V\RVTNRDI\ICQMLYA/RRIEGD MIVCAAYCTPNLPKIWV*RVGL TNY\AAAY\CTGLLL\ARRLLP*/ RFGMDKIYEGQV\ELTGDEYNV ESIDGQPGCLPPAYL\DAGPCSE PTHWPLKFFGAPEGKLWMGGL SIPSQYPNRFP\GYDSEKQRNFN AE\VHRKAHSWGQNCCRFTWR YF\MEED\EDAYKK\QFS\QYIK\ NSVT\PDMM\EEMY\KKAHAAI RENPVLWKRRPK\KEV\KKKRW NRPKMS\LAQ\KKDRVAQKKAS FLQSSRSGLLESLTPAIFP
17457	47825	Α	17563	2	4967	
17458	47826	A	17564	56	284	RQTRRAAGIVRGDCPASSPTNC DLGQC/NSTS/LEPRFVQTVHLL CPWT/GGSSPLLAARGSLPVSLL GPLKPMVSCAGR
17459	47827	Α	17565	1	3525	
17460	47828	В	17566	35	3691	
17461	47829	В	17567	80	3787	
17462	47830	Α	17568	225	507	
17463	47831	A	17569	2	2621	
17464	47832	Α	17570	14	1658	
17465	47833	Α	17571	1	3654	
17466	47834	В	17572	1	3570	
17467	47835	Α	17573	1	3663	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
17468	47836	A	17574	485	2347	TEPKTKTT*LSQ*MQKRPLTKF
						NNTSC*KLSIN/IVLEVLARAIRQ
						EKEIKGIQLGKEEVKVSLFADD
						MIVYLENPTVSAQNLLKLIGNF
						SKVSGYKINVQKSQAFLYTNNR
						QTERQIMSELPFTIASKRIKYLGI
						QLTRDVKDLFKENNKPLLKEV
						KEDTNEWKNIPCSWVGRINIVK
						MAILPKVIYRFNAIPIKLPMTFF
		1				TELEKTTLKFIWNQKRACIAKSI
		1				FSQKNKAGGITLPDFKLYYKAT
l		}				VTKTAWYWYQNRDIAQWNRT
		ļ				EPSEIMLHIYNYLIFDKPEKNKQ
						WGKDSLFNKWCWENWLAICR
						KVKLDPFLTPYTKMNSRWIKD
						LNVRPKTIKTLEENLGITIQDIG
						VGKDFMSKTPKAMATKAKIDK
				İ		WDLIKLKSFCTAKETTIRVNRQ
						PTTWEKIFATYSSDKGLISRIYN
						ELKQIYKKKTNNPIKKWAKDV
						NRHFSKEDIYAAKKHMKKCSS
						SLAIREMQIKTTMRYHLTPVRM
		ļ				AIIKKSGNNRKIQ/GGIWCDRIL*
						R*TTCRVAKEIQSL*RRI/WKRL
		1				QRTLSIPVLDAV*PPMF*ASVID
						TMTI*CFEARDTCFTLTLESFWD
1						MHRCLAASKGIGLLLC*PLIWH
	l	<u>L</u> .				MSLMGVKSPPFVFSCLWTSAV
17469	47837	В	17575	1	3354	
17470	47838	Α	17576	1	3297	

SEQ ID			SEQ ID NO:		1	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		0)/340,21/	sequence	or peptide sequence	deterion, (-possible nucleotide insertion)
17471	47839	Α	17577	1	2563	MKAEIKMFFETNENKDTTNQN
						LWDAFKAEEVESLNRPITGAEI
	ļ					GAIINSLPTKKSPGPDGFTAEFY
						QRYKEELVPFLLKLFQSIEKEEI
						LPNSFYEASIILIPKPGRDTTKKE
						NFRPISLMNIDAKILNKILANRI
						QQHIKKLIHHDQVGFFPGMQG
						WFNIRKSINVIQHINRAKDKNH
						MIISIDAEKAFDKIQQPFMLKTL
						NKLGIDGTYFKIIRAIYDKPTAN
						IILNGQKLEAFPLKTGTRQGCPL
						SPLLFNILLEVLARAIRQEKEIK
		İ				GIQLGKEEVKLSLFADDMIVYL
						ENPIVSAQNLLKLISNFSKVSGY
						KINVQKSQAFLYTSNRQTESQI
						MSELPFTIASKRIKYLGIQLTRD
]						VKDLFKENYKPLLKEIKEDTNK
		ĺ				WKNIPCSWVGRINIVKMAILPK
						VIYRFNAIPIKLPMTFFTELEKTT
						LKFIWNQKRAHITKSILSQKNK
						AGGITLPDFKLYYKATVTKTA
						WYCYQNRDIDQWNRTEPSEITP
						HTYNYLIFDKPEKNKQWGKDS
						LFNKWCWENWLAIWRKLKLD
						PFLTPYTKINSRWIKDLNVRPKT
						IKTLEENLGITIQDIGMGKDFMS
						RTPKAMATKAKIDKWDLIKLK
						SFCTAKETTIRVNRQPTTWEKIF
						ATYSSDKGLISRIYNELKQIYKK
						KTNNPIKKWEKDMNRHFSKED
						IYAAKKHMKKCSSSLAIREMQI
						KTTMRYHLTPVRMAIIKKSGNN
17472	47840	Α	17578	1	3514	MELKTKARELREECRSLRSRCD
						QLEERVSAMEDEMNEMKREG
						KFREKRIKRNEQSLQEIWDYVK
						RPNLRLIGVPESDVENGTKLEN
:						TLQDIIQENFPNLARQANIQIQEI
						QRTPQRYSLRRATPRHIIVRFTK
						VEMKEKMLRAAREKDRSTRQK
						VNKDTQELNSALHQADLIDIYR
1						TLHPKSTEYTFFSAPHHTYSKT
]						DHIVGSKALLSKCKRTEIITNYL
						SDHSAIKLELRIKNLTKSRSTTW
		ŀ				KLNNLLLNDYW
17473	47841	A	17579	1	3693	
	1.,,,,,,	۱٬۰	1.1017	L^	12222	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	•	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
17474	47842	A	17580	11577	3354	TEPKTKTT*LSQ*MQKRPLTKF
11/4/4	7 / 072	^	17300	1377	3334	NNLSC*KLSIN/IVLEVLARAIRQ
						EKEIKGIQLGKEEVKLSLFADD
						MIVYLENPIVSAQNLLKLISNFS
						KVSGYKINVQKSQAFLYTNNR
						QTESQIMSELPFTIASKRIKYLGI
						QLTRDVKDLFKENYKPLLKEIK
		i				EDTNKWKNIPCSWVGRINIVK
						MAILPKVIYRFNAIPIKLPMTFF
						TELGKTALKFIWNQKRARITKSI
						LSQKNKAGGITLPDFQLYCKAT
						VTKTAWFPSGDVGLEADFSPSH
		1				TLKTQFFSCLAEFAAASCFFQR
						MNGFGMAMTTTYSTGAAESPL
						PSCSIDQGDDTKLHRARSPGRT
						FPAAAGIPAAAAPDGPPLSLLH
			ŀ			KLWFPVELGGRALPRAEESHGE
		ŀ				VAALGVMVVAQGGKNQGEEA
						RSTPWLRPTSHLPPCSSSSAWW
						TEQTDAHPLLFLLCLGIYLLNA
-						LSNLSMVALVRSDGALRSPMY
						YFLGHLSLVDVCFTTVTVPRLL
						AGLLHPGQAISFQACFAEMYFF
		1				VALGITESYLPAAMSYDRATAA
		1				CRPLRYGALVTPWALRLAARY
						DRLASVVYAVITPTLNPFINSLR
						NKEVKGALKRGLRWRAAPQE
17475	47843	Α	17581	1	3042	
17476	47844	В	17582	1	3384	
17477	47845	В	17583	1	3264	
17478	47846	Α	17584	1	3780	
17479	47847	Α	17585	1	3165	
17480	47848	Α	17586	1	3894	
17481	47849	Α	17587	1	3345	
17482	47850	Α	17588	523	3981	
17483	47851	Α	17589	1	3429	
17484	47852	Α	17590	1	3921	
17485	47853	В	17591	1	4797	
17486	47854	Α	17592	1	6499	MVKGSIQQEELTILNIYAPNTG
						APRFIKQVLSDLQRDLDSHTLI
						MEDFNTPLSTLDRSTRQKVNK
						NTQELNSALHQADLIDIYRTLH
						PKSTEYTFFSAPHHTYSKIDHIV
						GSKALLSKCKRTEIITNYLSDHS
						AIKLELRIKNLTQSRSTTWKLN
						NLLLNDYWVHNEMKAEIKMFF
						ETNENKDTTYQNLWDAFKAVC
						RGKFIALNAYKRKQERSKIDTL
						TSQLKELEKQEQTHSKASRRQE
	<u> </u>					ITKIRAELKEIETQ

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	1	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
17487	47855	A	17593	1359	2520	IQCHPHQATNYFLHRIGKNYFK LHMEPNKSLHSQDNPKQKGTK LEV/CTLPDFKLYYKATVTKTA LYLYQ/NQRCRSMEQNRALRN NATYIYNYLIFDKPEKNKQWG KDTLFNKWCWENWLAICRKLK LDPFLTPYTKINSRWIKDLNVRP KTIKTLEENLGNTIQDIGMGKD FMSKTPKATATKAKIDKWDLIQ LKSFCTAKETTIRVNRQPTEWE KIFGIYSSDKGLISRIYNELQQIY KKKTNNPIKKWVKDMNRYFSK EDIYAAKRHMKKCSSSLAIREM QIKTTMRYYLTPLRMAIIKKFG NNRQETVREMACLLSGTMDFR HRACFVLSKPDFSTRWPGADA QRPPVLVVVALDSSLPFPQASG QRPPSLEDVDGGAGALVAPC
17488	47856	Α	17594	1	1227	
17489	47857	A	17595	2	2774	NLCSLIIPLREVTIVEKADSSSVL PSPLSISTRNRMTFLFANLKDRD FLVQRISDFLQQTTSKIYSDKEF AGSYNSSDDEVYSRPSSLVSSSP QRSTSSDADGERQFNLNGNSVP TATQTLMTMYRRSPEEFNPKL AKEFLKEQAWKIHFAEYGQGIC MYRTEKTRELVLKGIPESMRGE LWLLLSGAINEKATHPGYYEDL VEKSMGKYNLATEEIERDLHRS LPEHPAFQNEMGIAALRRVLTA YAFRNPNI
17490	47858	A	17596	1	3472	
17491	47859	Α	17597	1	900	
17492	47860	A	17598	152	1290	
17493	47861	A	17599	153	3834	
17494 17495	47862	A	17600 17601	299	1184 729	
17495	47863 47864	A A	17602	1	2232	
17497	47865	A	17603	36	224	TQWGPSVCWCLIPWPSFPDQ*P GF*AVLLPWRCQGFWPLLLAC APAGPATAAEPAVSSPAEP
17498	47866	Α	17604	1	624	
17499	47867	A	17605	1	578	RDLDLFSLPQDITL\VHYVDGIM LIGSSEQEVASTLDLLVRHLHA RGWEINLTKIQGLATSVKFLGV QWYGACRDIPSKVKDKLLHLA PPTTKKEAQHLAASFEWGPEQ KGLQQVQVAVQAALPLGLYDP ADPMVLEVSVADRDAVWSLW QAPIGESQWRPLGFQNKALPSA DNYSPFERQLLACYWALVETE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
		<u> </u>				
17500	47868	Α	17606	1	282	PIYRVSRKAANFEWSPEQEKAL
						QQVQAAVQAAWPLGPYDPAD
		1				PMVLEVSVADRDADWSCWQA
		1				SI/GHKVGHAQQHSIIKWKWYI
		<u> </u>				RDWARADPEGT
17501	47869	A	17607	1	338	SIYLVTQKAA/SSEWGPEQEKA
						LQEVQAAVQAALILEPYDPAGP
						VVLEVSLADRDAVWSLWQAPI
						GESQQRPLGFWSKALPSSA/DH KACHAQQHSIIKWKWYIHDRA
						RAGPEGT
17502	47870	A	17608	1	757	A COLOR
17503	47871	A	17609	1	1317	
17504	47872	Α	17610	2	24	LCQRLLLAEPNEKPGSLGNVM
}		}				AVARIEIGICEYYHEKTTEKALD
						SHGVLAGSTIKGVRSFQRNLEL
						KLPATERATANAIELLTVLDQA
						YENFAPQILPSTGSPTSQETAQF
						KANQNKPLVRGKGSPHEAIRYI
		1				SAAHREWKPAILTSAIRSFCST
						WLVFTSKNFPKLVTQHGSTIAG
		1				NGQSSDETQVQGAAWKSDSRG
						TKRQIPTWILAEGNNAGAQLDI
		1				PGPTIPAPNCSLKVPQSWSTTPS MPSSLGKAYWLLACYWALVET
						E/RLAMGHQVTM\KPELPVMN
						WVLSDPSSHKVGGAQQHSINK
		ŀ				WKWYIRNRARAGPEGTS*LPAP
						LIG
17505	47873	Α	17611	1	1815	
17506	47874	Α	17612	875	1916	PGKQGLEWTPRKLQQTCKRGA
						GQLEEKLANRKQ*HQH*QKG*P
						/PQKLNAKDHNSSPTREQNWTE
		}				NEFDELTEVGFRRWVITNSSEL
		1				KEHVLTQCKEAKNFDKRLEEL
						LIQITSLEKNISDLMELKNTARE/
						LLEAYTSINSQIDQAEMDR*PD*
		İ				*RRKERRIK*HNKK**RGYHN*S
		1				\QEIQTTIREYYKHLYANKLENL
		1				EEMDKFLDTYTLPRLNQEEVEP
						LNRPITSSEIEAVINSLPNQKSPG
						QTNSQLTSTRDAEKAFNKIQHP FMLKTLNKLDIDGTYLKITRAI/
						S*QTHSQCHTEWAKAGSIPFEN
						RHKTRMPSLTTPIQYSIGNSGQ
				1		GNQEFWKLHLVEREW
17507	47875	В	17613	1	1158	OTTO THE PROPERTY OF THE PROPE
1,501	1.,0,5		1.7015	T	L. 190	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
17508	47876	A	17614	851	1344	IILKPSPSPLYPWKNCLPRNPSL VPKRQTGSGVDIWQTAADLQQ RGLLEGKLTNRKE*YQHQQKR CPHQNPI*RSPTSNTKARE/QNW TENEFDELTEVGFRRWIITNSSQ LKEHLLTQCKEAKNLEKRLEEL LTRITSLEKNINDPMELKNTAEE PHEAYTSINS
17509	47877	В	17615	99	716	
17510	47878	Α	17616	1	588	
17511	47879	Α	17617	1	1140	
17512	47880	A	17618	1	774	
17513	47881	A	17619	3	527	AENSKKQNASSPPND/RNSS/EA QNWTENEFDELTEVGFRRWIV TNSSKLKEHVLTQCKEAKNLD KRLQELLTRITSLEKNINDLMEL KNTARELREAFTSFCVNCEAVS LITSLREDSHWLEDSKGHRM/R RHQE*KCEYFHPPHPGLQFYQL PESDSHSDPLWPHGGSFLSSVV
17514	47882	Α	17620	1	1383	
17515	47883	Α	17621	1	990	
17516	47884	В	17622	337	1515	
17517	47885	A	17623	153	474	QTKSNNVNIKKQDVHTKTPSEG HQHQRPK\DHNSLPAREQNWIE NEFDELTEVGFRRWVITNSSEL KEHVLTQCKEAKNLEKRLQEL LTRITSLEKNINDLMELKNTA
17518	47886	Α	17624			MPFSDTERVITGPSTKVHSYPG GLQALLLQLRLHSLMWRWSAA VAIGVHVRCDLLAVHPVLWAV PQWQEPIEKMLTLIPGPRITATII GLYQHEGIRAYCDSRVYYIKPS SGIESVLTRPSTQDHNSSPAREQ NWTENEFDELTEVGFKKWVIT NSFELKEHVLTQWKEAKNLEK RLANLLTRITSLENINDLMELK NTAQELHEAYASINSRINQSEER ISEIEDQLTEIEQVDKIREKRIK/R RNEQSL*EIWDYV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		
17519	47887	A	17625	53	814	PGKQGLEWTPRKLQQTCKRGA GQLEEKLANRKQ*HQH*QKG*P /PQKLNAKDHNSSPTREQNWTE NEFDELTEVGFRRWVITNSSEL KEHVLTQCKEAKNFDKRLEEL LIQITSLEKNISDLMELKNTARE/ LLEAYTSINSQIDQAEMDR*PD* *RRKERRIK*HNKK**RKEQAAI FAVLQPPLVITRQTGSGVDPQK TPADLQKRCRTVRRKTSQQKA VASTLTKRITTQKLNAKDHNSS PTREQNCTENEFDELTEVGFRR WVITNSSELKEHVLTQCKEAKN FDKRLEELLIQITSLEKNISDLM ELKNTARERWKHTQVSİAKSIK QKWTASQTNKEEKREESNSTIK NDKGDITTDPTEIQTTIREYYKH LYANKLENLEEMDKFLDTYTL
		1				PRLNQEEVEPLNRPITSSEIEAVI
						NSLPNQKSPGQTNSQLTSTR
17520	47888	A	17626	1	1566	NSEI NOKSI OQ INSQE1SIK
17521	47889	В	17627	1	1005	
17522	47890	A	17628	1	2172	
17523	47891	Α	17629	1	504	
17524	47892	Α	17630	2	1601	
17525	47893	С	17631	132	278	
17526	47894	A	17632	380	828	EVLRVMIELNAVIYPYPKDTGC QPLV*HAAASLNSPEVMGATQ LKPLRPCKKCPKI\LEGGH*EW MEARHRNLGTLHGLPCSRTHF* TTTAQGEWLCCLCHCGEHPQG GRHPCIC*HCYSCHTLAPPVQW TPNVAEPDNKAGAQYKSPRV
17527	47895	Α	17633	1	963	
17528	47896	A	17634	2	368	EFSDPIHHTFDHMWRTKEHNE AGWLLLSSVDKVMKENDELGD SISQLQKQILSLKSAKIALSESLI SCRERTEIVEK*T*ALIMLVADL Q*KVHAQPHHAQPRQVSTVKV RALIVLQEVTDP
17529	47897	Α	17635	1	1881	
17530	47898	A	17636	755	939	ARKNTRIIYSYGVCKNVETWR* MFCTLLGRQQGCATVGKGDRC GPLRYYASWRKGDVLQGD
17531	47899	Α	17637	367	423	
17532	47900	Α	17638	3	1925	
17533	47901	Α	17639	1	1137	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	i	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
17534	47902	A	17640	153	474	QTKSNNVNIKKQDVHTKTPSEG HQHQRPK\DHNSLPAREQNWIE NEFDELTEVGFRRWVITNSSEL KEHVLTQCKEAKNLEKRLQEL LTRITSLEKNINDLMELKNTA
17535	47903	Α	17641	1	1068	
17536	47904	В	17642	1	717	
17537	47905	A	17643	585	2302	VLQLLRQERLELFVPPGELVVS LASGVKLQTFAGGKAWLLQTH SDLGEQKLTTFIPISLSLKVIFWC NWPEAKTAAYDGEQATCEPVL KAMFLADTHLLREFLGHWLDK LRREWQVERAFQTALWLLQPE VVFILEDVFDEGKWSTSEAWV DDVEQFRKMFRHPSHVQLKVV AGNHDTGFHYETNTYEVERWR NRNNPSFIMGSITPTDYALSKCH LPPEDVVLIIYCGVKLPDRIVDS KAGLLKLPVLKNVISTQLQFPSI MDQLLVYNLFNETSPLVMCLD VMAMSNRGNHHLAATRLRKV LDHNSLPAREQNWTENEFDELT EVVFRRWVITSSSELKEC/DVLT QWKEAKNLEKRLDELLTRITSL QKNINDLIELKNTAQELHEAYT SINSQIDQVEERISEIEDQLNEIK CEDKIREKRMKRNKQTLQETW DYVKRPNPCWIGVPESDGENG TKLENTLQDII\QENFPNLARQA NIQIQEIQRTPQRYSSRRATPRHI IVRFTKVEMKEKMLRAAGEKE DETDNNEFFLWSQKGKRAGTL NDHVEESCPPTKNACLGTPIHC
17538	47906	A	17644	1	282	PIYRVSRKAANFEWSPEQEKAL QQVQAAVQAAWPLGPYDPAD PMVLEVSVADRDADWSCWQA SI/GHKVGHAQQHSIIKWKWYI RDWARADPEGT
17539	47907	Α	17645	3	903	
17540	47908	A	17646	1	349	
17541	47909	Α	17647	1	489	
17542	47910	Α	17648	1	352	
17543	47911	A	17649		505	KTDGSWRMTVDYHKLNQVAT PVAAAVPDVVSLLEQINTSLDT WYAAIDLGNVFFSIPVHKVHQK KFAFSWQGQQYTFTVLPQGYIN SPALCRHFVRRHLD/RFSLPQDI TVVHYIDDIMLVG\PLIYRMTR KAAIFEWGPEQKALQQVQAAV QAALPLGPYDPADPMRLR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	•	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	I .	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide inscrtion)
				sequence		
17544	47912	A	17650	1	1557	MASPEAVASQDNDGFSQKPPPT
						PLFASRSVNRLKSQRTPRGEVE
						SVTDGEKTDGSWRMTVDYCKL
						NQVVTPIAAAVPDVVSLLEQIN
-						TSPDITLVHYIDDIMLNGSSEQE
						VTNTLDLLVRHLCATGWEINPT
İ						KIQGSSTSVKFLGVQWCGDCR
						DVPSKVKDKLLHLAPPTTKKEA
						QHLTGLFGFRRKYIPYLGVLLC
		ŀ				PIYQVTRKAASFQWRPEQEKAL
						QQVQAAMQAALPLGPYDPAGP
						MVLEIAVADTEAVWGH/WHSA
İ						SFPSHPCHCPMGP*TKWP*WQG
						WRLCMGSATWTSTHEG*PGYG
						HQ*VPNLPAAETNTEPAI*HHS/
						SRRSASYLVAG*LYWTSCIMER
						AE\HLLWIWVCLSCMQCFCQD
					l i	YHPWTHGMPYPPLWYSTQLCL
						1
						*PRHSLYH*RSVAVGSCSWNSL
		ŀ				VLPCSPSS*SSRVDRTVERPFEIT
						TTMPTSQESRVQESRGGSGSGT
						THHHP**STSNIFASCFYGITFC
						WSRGLSSRGRNAATRRQNNSIK
17545	47913	В	17651	47	838	L*VKIATWTLWAPPTSKL
17546	47913	A	17652	1	1566	
17547	47915	A	17653	1	981	MVAIAKVLIAYKEDSDSHNQPN
11/34/	4/313	^	17033		761	TRMTVSKGPVSAVEGGRRGQD
İ						SDDCVEENDAVRTWPPSRILRT
						ESHPSPHTCSWRRHLLSSCPYT
						GMSFPIPSSGNISSTHAVISFPPL
						SEEMNPMLPKATVMTSLEAAA
1						RQNLKSRQAPRCEVESVTHEEV
İ						YYTRKELFEFSNLYRKKSGEQI
						REWILRKTDGSWRITVDYRKL
						NQMVTPIAAAVPDV/VVSLLEQ
						INTSPGTW\LCHNLIWRDLDCFL
						LPQNITLVHYTDDIMQIGSSEQE
						VANTLDLLPALMASW/ENSP*SI
						DRGRED*SLVHKWFCTICRHHP
L						KVDSCSTTAPF*DIPEGQQL
17548	47916	В	17654	179	1219	
17549	47917	Α	17655	1	2655	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
17550	47918	Α	17656	1	942	MVGKAKWKPLELPLPRKIVNQ
						KQHHIPEGIAEIAATIKDLKDAG
						VVIPTTSPFNSPIWPVQKTDGS
						WRMTVDYCKLNQVVTPIAAAV PDVVSFLEEINTSLGTWYAAID
		1				LANAFFSIPVHKVHQKPFAFSW
		1				QG/QQYTFTVLPQDYINSLAL*H
		1				NLIWRDLDYF\LLLQDITLVHYI
					·	DDIMLIGSNDHKVGGAQQHSII
						KWKLYIHDQAQTGPEGTTTSVI
						AQWAHEQSGPGSRDGGYAWA
						QQHGLPLTKADLATTTAECPVC
						QQQRPTLSPRYGTIPSLPLTKAL
						TLQLKKCSSGPMLMEFTGLAM
						FPIILKQLD
17551	47919	Α	17657	1	868	
17552	47920	Α	17658	83	346	
17553	47921	Α	17659	1	643	MVGKAKWKPLELPLPRKIVNQ
						KQHHIPEGIAEIAATIKDLKDAG
		l				VVIPTTSPFNSPIWPVQKTDGS
						WRMTVDYCKLNQVVTPIAAAV
						PDVVSFLEEINTSLGTWYAAID
	ŀ				LANAFFSIPVHKVHQKPFAFSW QG/QQYTFTVLPQDYINSLAL*H	
ļ						NLIWRDLDYF\LLLQDITLVHYI
	ļ					DDIMLIGSNDHKVGGAQQHSII
						KWKLYIHDQAQTGPEGT
17554	47922	c	17660	38	202	
17555	47923	Ā	17661	3	410	LKSRQAPRCEVESVTHEEVYYT
						RKELFEFSNLYRKKSGEQIREWI
	l	ı				LRKTDGSWRITVDYRKLNQMV
	l					TPIAAAVPDV/VVSLLEQINTSP
						GTW\LCHNLIWRDLDCFLLPQN
						ITLVHYTDDIMQIGSSEQEVAN
						TLDLL
17556	47924	Α	17662	2	526	
17557	47925	Α	17663	2	431	
17558	47926	В	17664	55	1506	
17559	47927	Α	17665	1	753	OL A PRODUMENT POLICY OF GERMAN
17560	47928	Α	17666	73	310	QLAEPPHWLPD*KGAGFEWGP
ł						EQKKALQ*VQAAVQAALPLGP
						YDPADPMVLVVSVADKDAVWI
17561	47929	A	17667	323	687	FHLSGSDRWRRTYRCL KTDGSWR/TVDYRKLNQVVTPI
1/301	41727	A	1 /00 /	1323	007	AAAVPDVVSLLEHINISPGIW/H
1						HQKQFAFS*QGQQFT/FTVL/PH
						GYINSLALCHNLIRRDLDHFSLP
						QDITLVHYIDDIVLTGSSEQEVA
1						GS\LESTCRHTKV
17562	47930	A	17668	3	1363	00.0001044444
17563	47931	A	17669	1	1203	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	I .	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
17564	47932	В	17670	130	1374	
17565	47933	Α	17671	1	1063	
17566	47934	Α	17672	3	239	
17567	47935	A	17673	231	368	NKHQEDSLKPPIPGPSSWTAF/P KPTLGQKGTCCTEKKKKKKK GRI
17568	47936	Α	17674	1	1270	
17569	47937	С	17675	100	270	
17570	47938	Α	17676	459	1978	
17571	47939	A	17677	35	347	VVRAWYPGWGQGSEKEWQW DQQEALGRVRGSKVTGEWEGR AGPEGTGQNGEGPRGILEVGGT TAPFCLAEGEWGPGGHPHHLLF PPSPPRPPPQPPPTPPEPPS*PPSP PRPPPQPPPTPPEPPS
17572	47940	A	17678	236	397	NWLQPCG*AHPTGSGFPAPKLC PASPGAPFSNE\ILSEPGPTLAPM PT/GAPCRSHGAGHSGDPGWPS SSPTGEDTNSAGPGAARVQMG CGHCVSLVPR*/CCWTPGGA
17573	47941	A	17679	132	492	TDHRDRKDKHPPALPASAMDK KNA/GQEGRPGASRGSSPMATA TTGPQGVMPGYH*CSLRPKGSS VGLW*MLPGLGLTLQDSGLPS GPRQVQNYCPGAKALNQGPQE PA*CKASNQRPSVYLPTREYPS EQIIVTEKTNILLRYLHQQWTK RTLPRRKTRSKQRKLTYGYCH YRSTGSNARLPLMFIKTQGFFS RLVVDAAWPGTHPSGQWAPF WPKAGPELLSRSQGFESGTSRA RLMFYPTMAAPVPK
17574	47942	Α	17680	618	755	
17575	47943	Α	17681	796	1092	
17576	47944	Α	17682	1	391	
17577	47945	A	17683	1	697	
17578	47946	Α	17684	1	888	
17579	47947	Α	17685	749	1667	
17580	47948	Α	17686	1	2088	
17581	47949	A	17687	189	438	
	47950	A	17688	387	497	_

NO: of peptide sequence hod in USSN location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
17583 47951 A 17689 D	i i	
117583 147951 IA 17689 11 1		
11,303 [1,731 [1]1,7007 [1	728	MAQEIEGDRLFRARLTRRVQG
		KGPCGPLKAPLGMVPECHSVK
		PSRDTFDSTKSFKALLLSFLKIP
		AIDVHKPIDHPINFTFPDPGHRA
		PARHRHSTAAAVTMSPSGP/PA
		AAPSPCSPIPRHG*GPLRATSYC
		PVSWKQKQNSVSNKQTNEQTK
		TAKAEGARSRGPRNKIPGLPPG
	1	PRSGLRIWLPPPREGSGFQTPAR
		GHPQPPCSGKGTECQSIAQNLA
		SRISPVGTSCGRTSRQTIRVRK
17584 47952 A 17690 I	819	ordor vorso orribrity rint rat
	2115	MSGSYSSVWAEDDIQFDSRFLE
		LKGDTKIDLKRFSSQGYVEPGK
		YNLQVQLNKQPLAEEYDIYWY
		AGEDDVSKSYACLTPELVAQF
		,
		GLKEDVAKNLQWSHDGKCLKP GQLEGVEIKADLSQSALVISLPQ
		AYLEYTWPDWDPPSRWDDGIS
		GIIADYSITAQTRHEENGGDDS
		NEISGNGTVGVNLGPWRMRAD
		WQTNYQHTRTQLALGEDYLNS
		DIFDGLNYVGGSVSTDDQMLPP
		HLRGYAPDISGVAHTTAK
	1080	
	253	
	1069	
	764	
	990	
1 . 1	1422	
	1017	
	431	
17594 47962 A 17700 1	846	
17595 47963 A 17701 I	1815	
17596 47964 A 17702 I	1578	
17597 47965 A 17703 1	1149	
17598 47966 A 17704 22	441	
17599 47967 A 17705 828	920	
<u> </u>	231	MESKRLDNA/CAGGGD*PQLH
		QCPR*TAVD*RRNQTAFA*RDA
		STYRHESGGNASPECHGLYQR
		QKNADGGGGQRRI
17601 47969 A 17707 1	563	
<u> </u>	372	
	491	
	969	
17605 47973 A 17711 59	839 725	
17000 47074 4 17710 100		ı
17607 47975 A 17713 1	251	
17607 47975 A 17713 1 2 17608 47976 A 17714 1381 2		

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	in USSN	location of first	codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
17610	47978	В	17716	1	2361	
17611	47979	Α	17717	1	2832	
17612	47980	Α	17718	1	3477	
17613	47981	В	17719	143	3122	
17614	47982	Α	17720	1	3018	
17615	47983	Α	17721	1	4371	
17616	47984	В	17722	1	2010	
17617	47985	Α	17723	1	2168	
17618	47986	Α	17724	1	2067	MKEIEIQKSLPKINESRSWFFEKI
		ŀ				HKIDRLLARLIKKKREKNQIDTI
						KNDKGDITANPTAIQTTIREYY
						KHLYANKLENLEEMDKFLDTY
	:				ļ	TLPRLNQEEAESLNTPITGSDIE
		İ				AIINSLPTKKSPGPDGLTAEFYO
		ŀ				RYKEELKKEGILLSSFYEASILLI
						PKPGRDTTKKENFRPISLMNID
		l				AKILNKILANRIQQHIKKLIHHD
						QVGFIPGMQGWFNICKSINVIQ
						HINRTKOKNHTIISIDAEKAFOK
				ĺ		IQQPFMLKTLNKLGIDGTYLKII
						RAIYDKPTANIILNGQKLEAFPL
		1				KTGTRQGCPVSPVVFNTVLEVL
		1				AREIRQEKEIKGIQLGKEEVKRS
						LFADDMIVYLENPIVSAQNLLK
						LISNFSKVSGYKINVQKSQAFL
						YTNNRQTESQIMSELPFTIAPKR
			ł			IKYLRIQLTRDVKDLFKENYKL
		l				LLNEIKEDTNKWKNIPCSWVGR
						INIVKMAILPKGIHRFNAIPIKLP
						MTFFTELEKTTLKFIWNQKRAR
						IAKSILSRKYKAGGSMLPDFKL
	ļ					CYKATVTKKAWYWYQNRDID
						QWNRTEPSEIMPHIYNYLIFDKP
						EKNKQWGKDSLFNKWCWENW
						LTICRKLKLDPFLTPYTEINSRW
	1		•			IKDLNVRPKTIKTLEENLGNTIQ
						DIGMGKDFMSKTPKAMATKAK
					1	IDEWDLLKLK\SFCTAKETTVR
			1			1
17610	47007	 	17705		2457	VNRQPTEWETIFAIYSS
17619	47987	A	17725	1	3457	
17620	47988	A	17726	1	3630	
17621	47989	В	17727	1	3384	
17622	47990	В	17728	1	2199	
17623	47991	A	17729	i .	3924	
17624	47992	Α	17730	1	3345	
17625	47993	Α	17731	1	2382	
17626	47994	Α	17732	1	3720	
17627	47995	A	17733	1	3894	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
17628	47996	A	17734	1	2427	MGCVRQIIPLGTHPVVHQLTSIP
						QNCQSPQKTRKDREIGQPRGAY
1						GDMRSKCNVMLMMESWNRKR
ŀ						TLEIQTTIREYYKHLYTNKLENL
ł						EEMDKFLDTYTLPRLNQEEVES
						LNRPITHSDIVTIINSLPTKK\SPG
		ŀ				PDRFTAKFYQRYKEELVPFLLK
1	-					LFQSIEK/EGILPNSFYEASIILIPK
}						LGRDTTKKENFRPISLMNTDAK
						ILNKILANRIQQHIKKLIQHDQV
ļ		ŀ				GFIPGMQGWFNIRKSINIIQHIN
		ŀ				RIKDKNHMIISIDAEKAFDKIQQ
İ						PFMLKTLNKLGIDGTYLKIIRAI
						YDKPTANIILNGQKLEAFPLKT
1						GTRQGCPLSPLLFNIVLEALAR
						AIRQEKEIKGIQLGKEEVKLSLF
						ADDMIVYLENPIVSAQNLLKLIS
						NFSKVSGYKINVQKSQAFLYTS
						NRQTESQIMSELPFTIASKRIKY
						LGIQLTRDVKDLFKENYKPLLK
					:	EIKEDTNKWKNIPCSWIGRINIV
						KMAILHKALYRFNAIPIKLPMT
			ļ			FFTELEKTTLKFIWNQKRACIA
						KSILSQKNKAGGITLPDFKLHY
		İ				KATVTKTAWYWYQNRDIDQW
ŀ						NRTEPSEIIPPIYNYLIFDKLDKN
						KKWGKDCLFNKWCWQNWLAI
						CRKLKLDPLLTPYTKINSRWIK
						DLNVRPKTIKTLEENLGITIQDI
ĺ						GMGKDFMSKTPKAMATKDKID
						KWDLIKLKSFCTAKETTIRVNR
						QPTKWEKIFVTYSSDKGLISRIY
17629	47997	Α	17735	1	3780	
17630	47998	Α	17736	1	2808	
17631	47999	A	17737	1	4842	
17632	48000	В	17738	331	3282	
17633	48001	Α	17739	1	2880	
17634	48002	Α	<u>17</u> 740	1	3213	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
17635	48003	A	17741		2561	MKAEIKTFETNKNKDTTYQNL WDTFKAVCRGKFIALNAHKRK QERSKIDTLTSQLKELEKQEQT YSKASRRQEITKIRAELKEIETQ KTLQKINESRSWFFEKINKIDRP LTRLIKKKREKNQIDSIKNDKG DITTDPTEIQTTIREYYKHLYAN KLENLEEMDKFLDTYTLPRLNQ EEVESLNRPITGSEIEAIINSLPT KKSPGPDGFTAEFYQRYKEELV PFLLKLFQSIEKEGILPNSFYEAS IILITKPGRHTTKKNFRPISLMNI GAKILNKILANRIQQHIKKLIHH DQVGFIPGMQGWFNIRKSINVI QHINITNDKNHMIISIDAENAFD KIQQRFMLKTLNKLGIDGMYL KIIRAIYDKPTATQNLLKLIGNF SKVSGYKINVQKSQAFLYTNNS QTESQIMTELPFTIASKRIKYLGI QLTRDVKDLFKENYKPLLSEIK EDTKKGKNIPCSWVGRINIMK MDILPKVIYTFNAIPIKLPMTFFT EVEKTTLKFICNQKRARIAKSFL SQKNKAGGITPPDFKLYYKATV TKTACYWYQNRDIDQWNRTEP SEIMQHIYNYLIFDKPDKNKQW GKDSLFNKWCWENWLAICRKL KLDSFLTPYTKINSRWIKDLNIR PKTIKTLEENLGSIIQDIGMGKD FMSKTPKAMATKAIIDEWDLIK LKSFCTAKETTMRVNRQPTEW
17636	48004	A	17742	1	5042	EKIFATYSSDIGLISRIYNELKQI MTGSNSHITILTLNINGLNSAIK RHRLASWIKSQDPSVCCIQETH LMCRDTHRLKIKGWRKIYQAN GKQKKAGVAILVSDKTDFKPT KIKRDKEGHYIMVKGSIQQEEL TILNIYAPNTGAPRFIKQVLSDL QRDLDSHTLIMGDFNTPLSILDR STRQKVNKDTQELNSALHQAD LIDIYRTLHPKSTEYTFFSAPHH TYSKIDHIVGSKALLSKCKRTEI ITNYLSDHSAIKLELRIKNLTQS
17637	48005	A	17743	101	471	RSTTWKLNNL MLSRILASRNRMRFAVEVVRA VRE/RVGNDFIIIYRLSMLDLVE DGGTFAETVELAQAVEAAGATI INTGIGWHEARIPTIATPVPRGA FSWVTRKLKGHVSLPLVTTNRI NDPQVADDILSSTH
17638	48006	Α	17744	161	311	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	I D	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
17639	48007	Α	17745	221	799	
17640	48008	Α	17746	20	346	
17641	48009	Α	17747	431	1578	
17642	48010	Α	17748	1	771	
17643	48011	A	17749	3	863	AVLRREMRQMEGAWLGSDLV DQGKERLNRLGFFETVDTDTQ RV/PGSPDQVDVVYK\IVSLGGR LFYNDFQADDADLSDYTNKKIE IIDSHADYFHIDIMDGHFVPNLT LSPFFVSQVKKLATKPLDCHLM VTRPQDYIAQLARAGADFITLH PETINGQAFRLIDEIRRHDMKV GLILNPETPVEAMKYYIHKADK ITVMTVDPGFAGQPFIPEMLDK LAELKAWREREGLEYEIEVDGS CNQATYEKLMAAGADVFIVGT SGLFNHAENIDEAWRVMTAQIL
17644	48012		17750	2	893	AAKSE
17645	48012	A A	17751	139	2057	
17646	48014	A	17752	2	361	
17647	48015	A	17753	497	897	PRSASFSAWLAFCST/TKDSHPL LAQLLDGVENLDHRQHLLFTA GHGARPLIRAHADVEENCDHPF ILIVHPSIHPIIYLSIHPSTIRGERL MTQIPLIKHFLAGVTGLKLTQQ MRFCENVCLPLIATAGNRQTH
17648	48016	A	17754	83	1038	PSRAGHLPPKIKSSMAGTTPCLS IMVRQRK/PLVPAPLYQGMRDG KIVRFEEITRTPLEVQDCLLASA LALGLALMGNAQAVTTIPFWH SMEGELGKEVDSLAQRFNAEN PDYKIVPTYKGNYEQNLSAGIA AFRTGNAPTYFAGHIMQACGV ALSATIYFIRRKVVSLSDQKLNI AMTFVGSMSGALLVQYVQAD VLRQILPILVICIGLYFLLMPKLG EEDRQRRMYGLPFALIAGGCV GFYDGFFGPAAGSFYALAFVTL CGFNLAKATAHAKLLNATSNIG GLLLFILGGKVIWATGFVMLVG QFLGAPHGVPTCV
17649	48017	Α	17755	571	1728	
17650	48018	Α	17756	332	511	
17651	48019	A	17757	581	637	RALR*SLQDGSAKRGASVS
17652	48020	Α	17758	1	1098	
17653	48021	A	17759	1	366	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
17654	48022	Α	17760	321	611	GEGENYCDSVVMTWVAI*QN
						WQCHCCSLRCASLRQG*RQDV
						TDTRGETIKREMARNWV/WRA
						RVINAGFSRGSLLRGPISDHRRK
						RLSASIRQFYPRP
17655	48023	В	17761	95	645	
17656	48024	В	17762	1	3978	
17657	48025	Α	17763	1	328	MLIAEGLMTAKNITVITNSLPA
1						AFALSENKDITLVVCGGTVRHK
1		1				TRSMHGSIAERSLQDINADLMF
]				VGADGIDAVNGITTFNEGYSIS
						GAMVTAANKVIAVLDSSKFNR
						RGFNQVLPIEKIDIHTDDAVSEV
						DKLALQKTRDTSLVYVLSLADF
		ŀ				FRTASTIGERDGTQVEMILFAGF
						VYLVISLSASLLVLTDCSTEVK KGEVVVVCGPSGSGKSTLIKTV
						NGLEPVQQGEITVDGIVVNDKK
						TDLAKLRSRVGMVFQHFELFPH LSIIENLTLAQVKVLKRDKAPA
						REKALKLLERVGLSAHANKFP
						AHYSGGQQQRVAIARALCMDP
						IAMLFDEPTSALDPEMINEVLD
						VMVELANEGMTMMVVTHEM
						GFAVKWRIGSTQQGAQLHCNR
						WLHVRKHLLVAYYNLVGIKPG
						KESYMRLNEKALDDFCQSLVD
						YLSAGHFSIYERILHKLEGNGQ
						LARAAKIWPQLEANTQQIMDY
		İ				YDSSLETAIDHDNYLEFQQVLS
		İ				DIGEALEAAFVLEDKLILLVLD
	-					AAR*NLLKLKVIIMIDGGFOTGI
						VVIHNLLSIGFELRPNLRGAREL
						PVAFQLMKNALINTEMSGRQVI
						DQALTKIIKGFFV
17658	48026	A	17764	216	473	DALRVASGNGCYSALLPLFCRL
			1			L*AESA*VPVQRPCGETLMRSS
						VVCLGGTAPECSSLGCAEKSGR
				1		SWPRKVAHTLGSWTVPRES
17659	48027	В	17765	1	1570	
17660	48028	A	17766	1580	1764	PASSLVMR*CLNRIA\SRAHQPA
						VRAY*SASAPGIA*KENA*G*RL
						HHR*NPAGKPDPGLL
17661	48029	Α	17767	1	1827	
17662	48030	В	17768	1	1281	
17663	48031	В	17769	72	1169	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	•	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
17664	48032	A	17770	20	552	SAGDA*DWFWHGGVAGGNG*
						RSP*A/VTSSALKCIHRALVRAW
			ļ			LLRMKKV*ATCA*CVTMRLKC
						CIK*/CPDNSLRMVQLFFPDPWH
						KARHNKRRIVQVPFAELVKSKL
ŀ						QLGGVFHMATDWEPYAEHML
						EVMSSIDGYKNLSESNDYVPRP
						ASRPVTKFEQRGHRLGHGVWD
						LMFERVK
17665	48033	Α	17771	3287	3568	CAFQRNIWHAKLNASPTPVFVP
		l				**LLASLTIPMKPAAMPGGKMD
		1				WWRVCRASASRPCQK*SLCQT
		1				PASVNTLLTVTAVCCASMASTT
						TRLWKI
17666	48034	A	17772	l	3126	
17667	48035	Α	17773	249	3943	
17668	48036	В	17774	1	2178	
17669	48037	Α	17775	263	434	
17670	48038	Α	17776	619	1184	MRTSCGSFIKRRVMRRIASGMV
						AENRAV**PSGICAIMVSTSSMK
						PMRSISSASSRTRPLSL/AEVQSA
						TFQVVQQTARSTDNDLRPLTQ
						GA*LHVITLAAVQS/IPHLRRAC
		1				V*RNPSLLQQPVPPVRGSVPAP
						ESVVLSAQDQDSATEVKRMLQ
		İ				FFRSPSGPYPERRGHSADVEYT
12671	40020	<u> </u>	17777	1	1405	LPELEMVFRNPDR
17671 17672	48039 48040	A A	17777 17778	497	1485 1838	
17673	48041	A	17779	534	922	
17674	48042	A	17780	1	458	MQQLEEALKQLAQGSGSSQAL
17074	10072	l^`	17700	'	150	TQVRRWDSACQKLPDANLALI
		i				SVAGEYAAELANQALDRNLNV
						MMFSDNVTLEDEIQLKTRARE
	į					KGLLVMGPDC/WYVDDCRHTA
						GFC*RDAGRQYWRHWRFRYR
						DSGAVFADCAGRGGNYSRDWP
						WRARPQP
17675	48043	A	17781	2	270	QAGKYYGGAVFAPVFGAIMGG
''''	""	`		_		VLRTMNIEPDALTSVDKNEFVI
						NHGEGTGGRS*FARPSCSVGAR
						RTFASTARDDTRQPCGCGGRSL
						CS
17676	48044	Α	17782	1	1437	
17677	48045	В	17783	63	201	
17678	48046	Α	17784	1016	1263	
17679	48047	Α	17785	1	933	
17680	48048	В	17786	1	1888	
17681	48049	Α	17787	845	2220	
17682	48050	Α	17788	110	581	
17683	48051	A	17789	696	989	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
17684	48052	Α	17790	199	573	
17685	48053	A	17791	1	3270	
17686	48054	Α	17792	1	3018	
17687	48055	В	17793	1	708	
17688	48056	A	17794	264	512	FLAQWLPELPLSRHLRQCTWCR RHRNRW*VRSLHRCWLADCRC PELRIRSHDL*SLAQRSLYDRTR TPDRRQLRTVLHWLLC
17689	48057	A	17795	1	2109	
17690	48058	A	17796	1239	1854	YHDDRNQPDAGNPSNADSGPQ I*QATGSPRRLVRR\RWKRVNR LLAGKLAIVIDDFGYRPHNENQ VLAMPSAISVAVLPDSPHAREM ATKAHNSGHEVLIHLPMAPLSK Q/TAGEKYATPGDEQRRN*AHY S*\WVNNVPYAVGINNHMGSK MTSNLFGMQKVMQALERYNL YFLDSVTIGNTQAMRAAQGTG VKVITRKVFLDGFAK
17691	48059	В	17797	1	2243	
17692	48060	A	17798	74	295	TISVLWLMLSLNTSKKRLA*TG SIGFFGLHSFTPRNGLRGASGTA FFGGVEVSTCGSFSKLLAFTSV MSGGRL
17693	48061	Α	17799	1	1872	MLRIHCMQHWYNLSDGAMED ALYEIASMRLFARLSLDSALPD RTTIMNFRHLLEQHQLARQLFK TINRWLAEAGVMMTQGTLVD ATIIEAPSSTKNKEQQRDPEMH QTKKGNQWHFGMKAHIGVDA KSGLTHSLVTTAANEHDLNQL GNLLHGEEQFVSADAGYQGAP QREELAEVDVDWLIAERPGKV RTLKQHPRKNRTGISIAYMTGSI RARVEHPFRIIKRQFGFVTARYR GCLNTITNGDVFPLATWFGG
17694	48062	Α	17800	1	1269	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ŀ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		İ		sequence		
17695	48063	A	17801	11	11988	MNDSEFHRLADQLWLTIEERLD
11000	10003	ľ.	1,,00,	<u> </u>	1,700	DWDGDSDIDCEINGGVLTITFE
						NGSKIIINRQEPLHQVWLATKQ
						GGYHFDLKGDEWICDRSGETF
						WDLLEQAATQQAGETITGFRV
						YTLQLNAQLRLGFLRLPYSVNL
						ATEYKSLTHYTKVPEATNPSLE
						i :
						SSVDVKLSANRLKYPMMRKRL
						MKMWREAKALHSDPVEAWAS
						IIEDADKAKSFKQARGRGGFVR
						SSWQEVNELIAASNVYTIKNYG
		į				PDRVAGFSPIPAMSMVSYASGA
						RYLSLIGGTCLSFYDWYCDLPP
					1	ASPQTWGEQTDVPESADWYNS
						SYIIAWGSNVPQTRTPDAHFFTE
						VRYKGTKTVAVTPDYAEIAKL
						CDLWLAPKQGTDAAMALAMG
						HVMLREFHLDNPSQYFTDYVR
						RYTDMPMLVMLEERDGYYAA
						GRMLRAADLVDALGQENNPE
						WKTVAFNTNGEMVAPNGSIGF
•						RWGEKGKWNLESIA\RIENPHF
					[RSVKHNPVLVRQLPVKNLTLV
						DGNTCPVVSVYDLVLANYGLD
						RGLEDENSAKDYAEIKPYTPA
						WGEQITRVPRQYIETIAREFADT
						AHKTHGRSMIILGAGVNHWYH
						MDMNYRGMINMLIFCGCVGQS
						GGGWAHYVGQEKLRPQTGWL
						PLAFALDWNRPPRQMNSTSFFY
						NHSSQWRYEKVSAQELLSPLA
17696	48064	Α	17802	3854	5726	
17697	48065	В	17803	1	1869	
17698	48066	Α	17804	391	1246	
17699	48067	Α	17805	58	1353	
17700	48068	Α	17806	872	2123	
17701	48069	Α	17807	1	1422	
17702	48070	Α	17808	217	509	
17703	48071	Α	17809	76	422	
17704	48072	Α	17810	2	288	
17705	48073	A	17811	1	1626	
17706	48074	Α	17812	779	832	
17707	48075	Α	17813	1	1818	
17708	48076	Α	17814	1	645	
17709	48077	A	17815	157	293	ALVCSSPWPSEKCKSKPQ*DTIS
						HQLEWQSLKSQETTGAGEDVE
						K
17710	48078	Α	17816	1	1446	
17711	48079	Α	17817	1	306	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
17712	48080	A	17818	71	513	TIRSGLGWSQMEMRNMLGTGA
						KKYKLPSENTINTSTQIN*KI*K
		ļ				KRINSSTHTLSQD*TRKKLNL*I
						DQ*QELKLWQ*SIAYQPKRVQD
						QMDSQ/REFYQRYKEELHINRA
		1				KDKNHMIISIDAEKAFDKIQQRF
						MLKTLNEVLMGRISK
17713	48081	A	17819	1	828	
17714	48082	В	17820	1	567	
17715	48083	Α	17821	1	1224	
17716	48084	Α	17822	1	897	
17717	48085	Α	17823	207	547	NHFLQGLGPLLQFPTSSSSPSET
						TSAWTLCPHHYQAFGQSHSTSL
						*EV/HKLSHIFMSSSEPSKLFQPL
1		1				PVTQFQSRFHIFRYLFSSIPLYW
						CQFTVLVHFPAADKDISVTTIP
17718	48086	В	17824	1	1128	
17719	48087	Α	17825	1	765	
17720	48088	Α	17826	1	952	
17721	48089	Α	17827	1570	2176	FLQSTHLTKG*YPESTMNSNKF
						TRKKQPNQRAKDMKRPFSKED
						IDAANKHMKKCSSSLAIREMQI
						KTTMRYHFTPVRMAIIKKSGNN
						RCWRGCGEIGTLLHCWWDCKL
						IQPLWKSMWRFLKDLELEIPFD
						PAIPLLGIYPKDYKSCCYKDTCT
						HMFIAALFTIAKTWNQPKCPTM
						IDWIKKMWHIYTMEYYGAIKR
						MSSCPL
17722	48090	Α	17828	1	531	
17723	48091	Α	17829	1	462	
17724	48092	Α	17830	876	1111	L L DOWNSON H L DI INDOGO CHIOLIN
17725	48093	Α	17831	2	276	LLPGFFCSNLAPVPPSGGHQHD
						LPYPSSA*NPGASGTGQARMGQ
:						SHQWAFLWRKDAHVRKESGY
						MYSLYDRAYGERTTGRRGRGR
17706	10001	_	17022		1200	GPIPRP
17726 17727	48094	A	17832 17833	1293	1290 1863	RSTPSMKE/QLATALPRHMTAE
1//2/	48095	Α	1/033	1293	1003	RMIRIANTEFRKVPALGNCDTM
					1	SFVSAIVQCPQLGLEPGSASGH
		1				1
		1				AYLLPFGNKNEKSGKKNVQLII
		1				GYRGMIDLARRSGQIASLSARV
		1				VREGDEFSFEFGLDEKLIHRPGE
		1		1	1	NEDAPVTHVYAVARLKDGGTQ
		1		1		FEVMTRKQIELVRSLSKADTD*
	10001	<u> </u>	1500	<u> </u>		PWMCPCPMKRKSAT
17728	48096	С	17834	11	786	

No.	SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
17729		-		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
17729 48097 A 17835 2 654		sequence		09/540,217		of peptide sequence	deletion, \=possible nucleotide insertion)
GTHRDNVLQVQSKLRADGCA EDYFSQSEQVLPSELAPDEAPC			l		sequence		
GTHRDNVLQVQSKLRADGCA EDYFSQSEQVLPSELAPDEAPC	17700	1,0007	1	17025		1054	ECD CD CCOVAND CD ETH A A ET
BEDYFSQSEQVLPSELAFDEAPC DVVNKYFNITLRTIMEGQSIVN VDEHRAARFFYQARQRDLYG AGGSNAICADVQHKFLRIGYR QAYPDAHIMMMSASLLQEGD VLVV HSGRISDVKAAVELAR KNGAKIICITHSYIISPIAKLADD IICSPAPETPLLGRNASARILQL LLDAFFVSVAQLNIEQANINMG KTGAM*KRP*NWQKRTGQRLJ LLDAFFVSVAQLNIEQANINMG KTGAM*KRP*NWQKRTGQRLJ VPINTIHE*RNWPILETAHQPR RRY*VVMPRQEYYN RRY*VVMPRQEYYN RRY*VVMPRQEYYN RRY*VVMPRQEYYN RRY*VVMPRQEYYN 17731 48100 B 17837 1 466 466 466 47733 48101 A 17839 1196 27322 48104 A 17839 1196 27322 48104 A 17840 B 1732 48104 A 17842 302 1716 48104 A 17842 302 1716 48104 A 17842 302 1716 48105 A 17843 249 909 48107 A 17845 902 1455 RRCVLRP/GLDIMRNRLNIRQQ QRLEQAAYEMTALRAATIELG LVRGLPHLRTIHRQLYQDIFDW AGGRRITLINHGFVEKSTTUVE NGPSDQSIAMRRPQTISSHHQ QKKLGJELTNCYSHSGLPIKN VTGILICRQTHARPARLARLIFI ANKTNTINALVDIEVDDFFTDM AGGRRITLINHGFVEKSTTUVE NGPSDQSIAMRRPQTISSHHQ QKKLGJELTNCYSHSGLPIKN VTGILICRQTHARPARLARLIFI ANKTNTINALVDIEVDDFFTDM AGGRRITLINHGFVEKSTTUVE NGPSDQSIAMRRPQTISSHHQ QKKLGJELTNCYSHSGLPIKN VTGILICRQTHARPARLARLIFI ANKTNTINALVDIEVDDFFTDM CQMCCGTGSSRJPFTEKLKRAAAG AGAILA WGTCASWGCYGAAP NPTQATPIDKVITDKPIKVPGC PIPDVMSAIITYMVTFDRLPDW RMGRPLMFYGQKIIHOKCYRR HFDAGEFYQSWDDDAAPKVT ACTKWAAKGLPPITEVPHTLE WVWSRLLAGGNGLDRGRFSG DSSWISYLSSAISVTMAPATTLI QTSKAALACAISTLCNGWPA VTLAQHRWMKWSGLAS	17729	48097	A	17835	2	654	1
DVVNKYPNITI,RTIMBGQSIVW VDEHRAARFFYQARQRDLYG AGGSNAICADVQHKFLRIGVR QAYPDAHIMMMSASLLQEGD VLVVTHSGRTSDVKAAVELAK KNGAKIICITHSYHSPIAKLAD IICSPAPETPLLGRNASARILQL' LLDAFFVSVAQLNIEQANINM KTGAM*KRP*NWQKKTGGRLI V*PIATIHR*RNWPILFAHQPR RRY*VVMPRQEYYN 17730 48098 A 17836 1 4806 48]		l				1
VDEHRAARFYQARQRDLVG							1
AGGSNAICADVQHKFLRIGYR QAYPDAHIMMMSALIQUE	}						1
QAYPDAHIMMMSASLLQEGD VLVYTHSGRTSDVKAAVELADN IICSPAPETPLLGRNASARILQL/ LLDAFFVSVAQLNIEQANINK KTGAM*KRPNWQKRTGQRLI V*PIATHR*RNWPILIFAHQPR RRY*VVMPRQEYYN RRY*VVMPRQEYYN 17730	1						•
VLVVTHSGRTSDVKAAVELAK KNGAKIICITHSYIISPIAKLAS IICSPAPETPILIGRASARILQL' IICSPAPETPILIGRASARILQL' IICSPAPETPILIGRASARILQL' IICSPAPETPILIGRASARILQL' V*PIATHIR*RNWPILIFAHQPRI RRY*VVMPRQEYYN V*PIATHIR*RNWPILIFAHQPRI RRY*VVMPRQEYYN V*PIATHIR*RNWPILIFAHQPRI RRY*VVMPRQEYYN V*PIATHIR*RNWPILIFAHQPRI RRY*VVMPRQEYYN V*PIATHIR*RNWPILIFAHQPRI V*PIATHIR*RNWPILIFAHQPRI V*PIATHIR*RNWPILIFAHQPRI V*PIATHIR*RNWPILIFAHQPRI V*PIATHIR*RNWPILIFAHQPRI V*PIATHIR*RNWPILIFAHQPRI V*PIATHIR*RNWPILIFAHQPRI V*PIATHIR*RNWPILIFAHQPRI V*PIATHIR*RNWPILIFAHQPRI V*PIATHIR*RNWPILIFAHQPRI V*PIATHIR*RNWPILIFAHQPRI V*PIATHIR*RNWPILIFAHQPRI V*PIATHIR*RNWPILIFAHQPRI V*PIATHIR*RNWPILIFAHQPRI V*PIATHIR*RNWPIATHIR*RN							· · · · · · · · · · · · · · · · · · ·
RNGAKIICITHSYIISPIAKLAD\ IICSPAPETPLLGRNASARILQ\ LLDAFFVSVAQLNIEQANINM\ KTGAM*KRP*NWQKRTGQRLI\ V*PIATIHR*RNWPIILFAHQPR\ RRY*VVMPRQEYYN 17730							
IICSPAPETPLLGRNASARILQL' LLDAFFVSVQLNIEQANINM KTGAM*KRP*NWQKRTGQRLI V*PIATIHR*RNWPILIFAHQPR RRY*VVMPRQEYYN 17730							i l
LLDAFFVSVAQLNIEQANINMC KTGAM*KRP*NVQKRTGQRLI							i e
RTGAM*KRP*NWQKRTGQRLI V*PIATIHR*RNWPIILFAHQPRI RRY*VVMPRQEYYN							1
17730							, , , , ,
17730		•	1				1
17730							V*PIATIHR*RNWPIILFAHQPRK
17731							RRY*VVMPRQEYYN
17732	17730	48098	Α	17836	1	4806	
17733	17731	48099	Α	17837	1	466	
17734	17732	48100	В	17838	193	1045	
17735	17733	48101	Α	17839	1196	2732	
17736	17734	48102	В	17840	1	732	
17737	17735	48103	Α	17841	290	664	
17738	17736	48104	A	17842	302	1716	
17739 48107 A 17845 902 1455 RRCVLRP/GLDIMRNRLNIRQQ QRLEQAAYEMTALRAATIELG LVRGLPHLRTIHRQLYQDIFDW AGQRRITLINHGFVPKSTTVVE NGPSDGSIAMRRPGQTISRHU QKKLGIELTNCYEHSGLPDIKN VTGILICRQTHARPARLARLIFI ANKTNTINALVDIEVDDFFTDA CQIDCCFETGSHSVLARLGGAS AQDLQLDATSASHGSK 17740 48108 A 17846 I 1668 17741 48109 A 17847 I 1080 17742 48110 A 17848 I 907 MRAGFTSNVLTFLTCTSHFLNT FRAADVANVQTATSNFRQIQV EDIITQYNGKYILAVEGNPPLG QGMFCISSGRPFIEKLKRAAAG ASAIIAWGTCASWGCVQAARP NPTQATPIDKVITDKPIIKVPGC PIPDVMSAIITYMVTFDRLPDV, RMGRPLMFYGQRIHDKCYRRA HFDAGEFVQSWDDDAA\PKVT ACTK WAAKGLPPITPVPHTLE* WVWSRLLAGGNGLDRGRFSG DSSWISYLSSAISVTMAPATTLI QTSKAALAACAISTLCNGWPA VILAQHRWMKWSGLAS	17737	48105	Α	17843	249	909	
QRLEQAAYEMTALRAATIELG LVRGLPHLRTIHRQLYQDIFDW AGQRRITLINHGFVPKSTTVVE NGPSDQSIAMRRPGQTISRIHQ' QKKLGIELTNCYEHSGLPDIKN VTGILICRQTHARPARLARLIFTI ANKTNTINALVDIEVDDFFTDA CQIDCCFETGSHSVLARLGGAS AQDLQLDATSASHGSK 17740 48108 A 17846 I 1668 17741 48109 A 17847 I 1080 17742 48110 A 17848 I 907 MRAGFTSNVLTFLTCTSHFLNT FRAADVANVQTATSNFRQIQV EDIITQYNGKYILAVEGNPPLGI QGMFCISSGRPFIEKLKRAAAG ASAIIAWGTCASWGCVQAARP NPTQATPIDKVITDKPIIKVPGC PIPDVMSAIITYMVTFDRLPDVI RMGRPLMFYQQRIHDKCYRRA HFDAGEFVQSWDDDAA\PKVT ACTKWAAKGLPPITPVPHTLE* WVWSRLLAGGNGLDRGFSGS DSSWISYLSSAISVTMAPATTLI QTSKAALAACAISTLCNGWPA VILAQHRWMKWSGLAS	17738	48106	Α	17844	36	340	
LVRGLPHLRTIHRQLYQDIFDW AGQRRITLINHGFVPKSTTVVE NGPSDQSIAMRRPGQTISRIHQ¹ QKKLGIELTNCYEHSGLPDIKN VTGILICRQTHARPARLARLIF1 ANKTNTINALVDIEVDDFFTDA CQIDCCFETGSHSVLARLGGAS AQDLQLDATSASHGSK 17740 48108 A 17846 l 1668 17741 48109 A 17847 l 1080 17742 48110 A 17848 l 907 MRAGFTSNVLTFLTCTSHFLNT FRAADVANVQTATSNFRQIQV EDIITQYNGKYILAVEGNPPLGI QGMFCISSGRPFIEKLKRAAAG ASAIIAWGTCASWGCVQAARP NPTQATPIDKVITDKPIIKVPGC PIPDVMSAIITYMVTFDRLPDVI RMGRPLMFYGQRIHDKCYRRA HFDAGEFVQSWDDDAA\PKVT ACTKWAAKGLPPITPVPHTLE* WVWSRLLAGGNGLDRGRFSG DSSWISYLSSAISVTMAPATTLI QTSKAALAACAISTLCNGWPA VILAQHRWMKWSGLAS	17739	48107	Α	17845	902	1455	RRCVLRP/GLDIMRNRLNIRQQ
AGQRRITLINHGFVPKSTTVVE NGPSDQSIAMRRPGQTISRIHQ' QKKLGIELTNCYEHSGLPDIKN VTGILICRQTHARPARLARLIFT ANKTNTINALVDIEVDDFFTDA CQIDCCFETGSHSVLARLGGAS AQDLQLDATSASHGSK 17740 48108 A 17846 1 1668 17741 48109 A 17847 1 1080 17742 48110 A 17848 1 907 MRAGFTSNVLTFLTCTSHFLNT FRAADVANVQTATSNFRQIQV EDIITQYNGKYILAVEGNPPLG! QGMFCISSGRPFIEKLKRAAAG ASAIIAWGTCASWGCVQAARP NPTQATPIDKVITDKPIIKVPGC PIPDVMSAIITYMVTFDRLPDV! RMGRPLMFYGQRIHDKCYRRA HFDAGEFVQSWDDDAA\PKVT ACTKWAAKGLPPITPVPHTLE* WVWSRLLAGGNGLDRGFSG DSSWISYLSSAISVTMAPATTLI QTSKAALAACAISTLCNGWPA VILAQHRWMKWSGLAS							QRLEQAAYEMTALRAATIELGP
NGPSDQSIAMRRPGQTISRIHQ QKKLGIELTNCYEHSGLPDIKN VTGILICRQTHARPARLARLIFT ANKTNTINALVDIEVDDFFTDA CQIDCCFETGSHSVLARLGGAS AQDLQLDATSASHGSK 17740 48108 A 17846 1 1668 17741 48109 A 17847 1 1080 17742 48110 A 17848 1 907 MRAGFTSNVLTFLTCTSHFLNT FRAADVANVQTATSNFRQIQV EDIITQYNGKYILA VEGNPPLG QGMFCISSGRPFIEKLKRAAAG ASAIIAWGTCASWGCVQAARP NPTQATPIDKVITDKPIKVPGC PIPDVMSAIITYMVTFDRLPDVI RMGRPLMFYGQRIHDKCYRRA HFDAGEFVQSWDDDAA\PKVT ACTKWAAKGLPPITPVPHTLE* WVWSRLLAGGNGLDRGFSG DSSWISYLSSAISVTMAPATTLI QTSKAALAACAISTLCNGWPA VILAQHRWMKWSGLAS							LVRGLPHLRTIHRQLYQDIFDW
QKKLGIELTNCYEHSGLPDIKN VTGILICRQTHARPARLARLIFT ANKTNTINALVDIEVDDFFTDA CQIDCCFETGSHSVLARLGGAS AQDLQLDATSASHGSK 17740 48108 A 17846 I 1668 17741 48109 A 17847 I 1080 17742 48110 A 17848 I 907 MRAGFTSNVLTFLTCTSHFLNT FRAADVANVQTATSNFRQIQV EDIITQYNGKYILAVEGNPPLG; QGMFCISSGRPFIEKLKRAAAG ASAIIAWGTCASWGCVQAARP NPTQATPIDKVITDKPIIKVPGC PIPDVMSAIITYMVTFDRLPDV; RMGRPLMFYGQRIHDKCYRRA HFDAGEFVQSWDDDAA\PKVT ACTKWAAKGLPPITPVPHTLE* WVWSRLLAGGNGLDRGFSG; DSSWISYLSSAISVTMAPATTLI QTSKAALAACAISTLCNGWPA VILAQHRWMKWSGLAS					}		AGQRRITLINHGFVPKSTTVVET
QKKLGIELTNCYEHSGLPDIKN VTGILICRQTHARPARLARLIFT ANKTNTINALVDIEVDDFFTDA CQIDCCFETGSHSVLARLGGAS AQDLQLDATSASHGSK 17740 48108 A 17846 I 1668 17741 48109 A 17847 I 1080 17742 48110 A 17848 I 907 MRAGFTSNVLTFLTCTSHFLNT FRAADVANVQTATSNFRQIQV EDIITQYNGKYILAVEGNPPLG; QGMFCISSGRPFIEKLKRAAAG ASAIIAWGTCASWGCVQAARP NPTQATPIDKVITDKPIIKVPGC PIPDVMSAIITYMVTFDRLPDV; RMGRPLMFYGQRIHDKCYRRA HFDAGEFVQSWDDDAA\PKVT ACTKWAAKGLPPITPVPHTLE* WVWSRLLAGGNGLDRGFSG; DSSWISYLSSAISVTMAPATTLI QTSKAALAACAISTLCNGWPA VILAQHRWMKWSGLAS				•			NGPSDQSIAMRRPGQTISRIHQV
VTGILICRQTHARPARLARLIFT ANKTNTINALVDIEVDDFFTDA CQIDCCFETGSHSVLARLGGAS AQDLQLDATSASHGSK 17740 48108 A 17846 I 1668 17741 48109 A 17847 I 1080 17742 48110 A 17848 I 907 MRAGFTSNVLTFLTCTSHFLNT FRAADVANVQTATSNFRQIQV EDIITQYNGKYILAVEGNPPLG; QGMFCISSGRPFIEKLKRAAAG ASAIIAWGTCASWGCVQAARP NPTQATPIDKVITDKPIIKVPGC PIPDVMSAIITYMVTFDRLPDV; RMGRPLMFYGQRIHDKCYRRA HFDAGEFVQSWDDDAA\PKVT ACTK WAAKGLPPITPVPHTLE* WVWSRLLAGGNGLDRGFFSG; DSSWISYLSSAISVTMAPATTLI QTSKAALAACAISTLCNGWPA VILAQHRWMKWSGLAS				[
ANKTNTINALVDIEVDDFFTDA CQIDCCFETGSHSVLARLGGAS AQDLQLDATSASHGSK 17740 48108 A 17846 1 1668 17741 48109 A 17847 1 1080 17742 48110 A 17848 1 907 MRAGFTSNVLTFLTCTSHFLNT FRAADVANVQTATSNFRQIQV EDIITQYNGKYILAVEGNPPLGI QGMFCISSGRPFIEKLKRAAAG ASAIIAWGTCASWGCVQAARP NPTQATPIDKVITDKPIIKVPGC PIPDVMSAIITYMVTFDRLPDVI RMGRPLMFYQQRIHDKCYRRA HFDAGEFVQSWDDDAA\PKVT ACTKWAAKGLPPITPVPHTLE* WVWSRLLAGGNGLDRGRFSG DSSWISYLSSAISVTMAPATTLI QTSKAALAACAISTLCNGWPA VILAQHRWMKWSGLAS							-
CQIDCCFETGSHSVLARLGGAS AQDLQLDATSASHGSK 17740 48108 A 17846 I 1668 17741 48109 A 17847 I 1080 17742 48110 A 17848 I 907 MRAGFTSNVLTFLTCTSHFLNT FRAADVANVQTATSNFRQIQV EDIITQYNGKYILAVEGNPPLGI QGMFCISSGRPFIEKLKRAAAG ASAIIAWGTCASWGCVQAARP NPTQATPIDKVITDKPIIKVPGC PIPDVMSAIITYMVTFDRLPDVI RMGRPLMFYGQRIHDKCYRRA HFDAGEFVQSWDDDAA\PKVT ACTKWAAKGLPPITPVPHTLE* WVWSRLLAGGNGLDRGRFSG DSSWISYLSSAISVTMAPATTLI QTSKAALAACAISTLCNGWPA VILAQHRWMKWSGLAS							N A
AQDLQLDATSASHGSK 17740 48108 A 17846 I 1668 17741 48109 A 17847 I 1080 17742 48110 A 17848 I 907 MRAGFTSNVLTFLTCTSHFLNT FRAADVANVQTATSNFRQIQV EDIITQYNGKYILAVEGNPPLG QGMFCISSGRPFIEKLKRAAAG ASAIIAWGTCASWGCVQAARP NPTQATPIDKVITDKPIIKVPGC PIPDVMSAIITYMVTFDRLPDV RMGRPLMFYGQRIHDKCYRRAHFDAGEFVQSWDDDAA\PKVT ACTKWAAKGLPPITPVPHTLE* WVWSRLLAGGNGLDRGFFSG DSSWISYLSSAISVTMAPATTLI QTSKAALAACAISTLCNGWPA VILAQHRWMKWSGLAS							
17740							1
17741 48109 A 17847 I 1080 17742 48110 A 17848 I 907 MRAGFTSNVLTFLTCTSHFLNT FRAADVANVQTATSNFRQIQV EDIITQYNGKYILAVEGNPPLGI QGMFCISSGRPFIEKLKRAAAG ASAIIAWGTCASWGCVQAARP NPTQATPIDKVITDKPIIKVPGC PIPDVMSAIITYMVTFDRLPDVI RMGRPLMFYGQRIHDKCYRRAHFDAGEFVQSWDDDAA\PKVT ACTKWAAKGLPPITPVPHTLE* WVWSRLLAGGNGLDRGRFSG DSSWISYLSSAISVTMAPATTLI QTSKAALAACAISTLCNGWPA VILAQHRWMKWSGLAS	17740	48108	Α	17846	1	1668	110000000000000000000000000000000000000
17742 48110 A 17848 I 907 MRAGFTSNVLTFLTCTSHFLNT FRAADVANVQTATSNFRQIQV EDIITQYNGKYILAVEGNPPLGI QGMFCISSGRPFIEKLKRAAAG ASAIIAWGTCASWGCVQAARP NPTQATPIDKVITDKPIIKVPGC PIPDVMSAIITYMVTFDRLPDVI RMGRPLMFYGQRIHDKCYRRA HFDAGEFVQSWDDDAA\PKVT ACTKWAAKGLPPITPVPHTLE* WVWSRLLAGGNGLDRGRFSGI DSSWISYLSSAISVTMAPATTLI QTSKAALAACAISTLCNGWPA VILAQHRWMKWSGLAS		!	-		ļ		
FRAADVANVQTATSNFRQIQV EDIITQYNGKYILAVEGNPPLGI QGMFCISSGRPFIEKLKRAAAG ASAIIAWGTCASWGCVQAARP NPTQATPIDKVITDKPIIKVPGC PIPDVMSAIITYMVTFDRLPDVI RMGRPLMFYGQRIHDKCYRRA HFDAGEFVQSWDDDAA\PKVT ACTKWAAKGLPPITPVPHTLE* WVWSRLLAGGNGLDRGRFSGI DSSWISYLSSAISVTMAPATTLI QTSKAALAACAISTLCNGWPA VILAQHRWMKWSGLAS					ļ		MRAGETSNYLTELTCTSHFLNT
EDIITQYNGKYILAVEGNPPLGI QGMFCISSGRPFIEKLKRAAAG ASAIIAWGTCASWGCVQAARP NPTQATPIDKVITDKPIIKVPGC PIPDVMSAIITYMVTFDRLPDVI RMGRPLMFYGQRIHDKCYRRA HFDAGEFVQSWDDDAA\PKVT ACTKWAAKGLPPITPVPHTLE* WVWSRLLAGGNGLDRGRFSGI DSSWISYLSSAISVTMAPATTLI QTSKAALAACAISTLCNGWPA VILAQHRWMKWSGLAS	1,,,,,,	10110	[]	17,570	1		
QGMFCISSGRPFIEKLKRAAAG ASAIIAWGTCASWGCVQAARP NPTQATPIDKVITDKPIIKVPGC PIPDVMSAIITYMVTFDRLPDVI RMGRPLMFYGQRIHDKCYRRA HFDAGEFVQSWDDDAA\PKVT ACTKWAAKGLPPITPVPHTLE* WVWSRLLAGGNGLDRGRFSG* DSSWISYLSSAISVTMAPATTLI QTSKAALAACAISTLCNGWPA VILAQHRWMKWSGLAS							
ASAIIAWGTCASWGCVQAARP NPTQATPIDKVITDKPIIKVPGC PIPDVMSAIITYMVTFDRLPDVI RMGRPLMFYGQRIHDKCYRRA HFDAGEFVQSWDDDAA\PKVT ACTKWAAKGLPPITPVPHTLE* WVWSRLLAGGNGLDRGRFSG* DSSWISYLSSAISVTMAPATTLI QTSKAALAACAISTLCNGWPA VILAQHRWMKWSGLAS							-
NPTQATPIDKVITDKPIIKVPGC PIPDVMSAIITYMVTFDRLPDVI RMGRPLMFYGQRIHDKCYRRA HFDAGEFVQSWDDDAA\PKVT ACTKWAAKGLPPITPVPHTLE* WVWSRLLAGGNGLDRGRFSG* DSSWISYLSSAISVTMAPATTLI QTSKAALAACAISTLCNGWPA VILAQHRWMKWSGLAS							-
PIPDVMSAIITYMVTFDRLPDVJ RMGRPLMFYGQRIHDKCYRRA HFDAGEFVQSWDDDAA\PKVT ACTKWAAKGLPPITPVPHTLE* WVWSRLLAGGNGLDRGRFSG* DSSWISYLSSAISVTMAPATTLI QTSKAALAACAISTLCNGWPA VILAQHRWMKWSGLAS							
RMGRPLMFYGQRIHDKCYRRA HFDAGEFVQSWDDDAA\PKVT ACTKWAAKGLPPITPVPHTLE* WVWSRLLAGGNGLDRGRFSG* DSSWISYLSSAISVTMAPATTLI QTSKAALAACAISTLCNGWPA VILAQHRWMKWSGLAS							·
HFDAGEFVQSWDDDAA\PKVT ACTKWAAKGLPPITPVPHTLE* WVWSRLLAGGNGLDRGRFSG* DSSWISYLSSAISVTMAPATTLI QTSKAALAACAISTLCNGWPA VILAQHRWMKWSGLAS							
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WVWSRLLAGGNGLDRGRFSG' DSSWISYLSSAISVTMAPATTLI QTSKAALAACAISTLCNGWPA VILAQHRWMKWSGLAS							· · · · · · · · · · · · · · · · · · ·
DSSWISYLSSAISVTMAPATTLI QTSKAALAACAISTLCNGWPA VILAQHRWMKWSGLAS	1			Ì	1		1
QTSKAALAACAISTLCNGWPA VILAQHRWMKWSGLAS					}		WVWSRLLAGGNGLDRGRFSGV
VILAQHRWMKWSGLAS							DSSWISYLSSAISVTMAPATTLN
			1				QTSKAALAACAISTLCNGWPA
17743 48111 A 17849 239 899		<u> </u>		<u> </u>			VILAQHRWMKWSGLAS
<u></u>	17743	48111	A	17849	239	899	

			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
17744	48112	Α	17850	155	249	CMRRKTALTGGEPDR*TAMR*
						HNKSPALTGS
17745	48113	Α	17851	1	1326	
17746	48114	Α	17852	1	1394	
17747	48115	Α	17853	46	705	
17748	48116	A	17854	2	488	
17749	48117	Α	17855	563	687	LLARTSRQ*KTSLNCVNKL*MP SGQWAVLLLPVCRCWWLRT
	48118	Α	17856	1	1429	
17751	48119	A	17857	1	1037	
17752	48120	A	17858	2	276	LLPGFFCSNLAPVPPSGGHQHD LPYPSSA*NPGASGTGQARMGQ SHQWAFLWRKDAHVRKESGY MYSLYDRAYGERTTGRRGRGR GPIPRP
17753	48121	A	17859	1	1290	
17754	48122	A	17860	1293	1863	RSTPSMKE/QLATALPRHMTAE RMIRIANTEFRKVPALGNCDTM SFVSAIVQCPQLGLEPGSASGH AYLLPFGNKNEKSGKKNVQLII GYRGMIDLARRSGQIASLSARV VREGDEFSFEFGLDEKLIHRPGE NEDAPVTHVYAVARLKDGGTQ FEVMTRKQIELVRSLSKADTD* PWMCPCPMKRKSAT
17755	48123	A	17861	207	210	QLSLLLNLPLDSIHHRLASSALK RVNDQVGAAPPRHQSPPPRNA VSLAPRPVPVEWSCAKLLCLSC RVNCQD*GNKALTSIHFVWGE GIVGKMWLFYCTEAHFVRDPS GYSN
17756	48124	A	17862	2	654	FCRQRQCQVYRRGRFTHAAFT GTHRDNVLQVQSKLRADGCAL EDYFSQSEQVLPSELAFDEAPQ DVVNKVFNITLRTIMEGQSIVN VDEIHRAARFFYQARQRDLYG AGGSNAICADVQHKFLRIGVRC QAYPDAHIMMMSASLLQEGDV VLVVTHSGRTSDVKAAVELAK KNGAKIICITHSYHSPIAKLADY IICSPAPETPLLGRNASARILQLT LLDAFFVSVAQLNIEQANINMQ KTGAM*KRP*NWQKRTGQRLF V*PIATIHR*RNWPIILFAHQPRK RRY*VVMPRQEYYN
17757	48125	Α	17863	1	4806	MAT V VIVIENCET IN
	48126	A	17864	1	466	
	48127	A	17865	214	907	
_	48128	A	17866	1288	2757	
17761	48129	A	17867	705	839	AGRSRENRRYRAKQPRCGKR*F *CVYRFYPSGRYAEPSSAPNYT

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
17762	48130	A	17868	199	1628	
17763	48131	A	17869	746	1013	TRARREITALRESSLVSASPCC
			1,,005			NWAAAFSS\YFCQQIVR*PHFDR
						QEKRSVQNQPDGCSSSASQSVR
		l				QRPKSPQQPVQSANPSLLLCR
17764	48132	A	17870	2	331	
17765	48133	Α	17871	1	591	
17766	48134	Α	17872	1	1197	
17767	48135	Α	17873	45	3402	NRFSHISVTSACGWYHIPVEE/Q
						PEY/PGNLELERRIRSAIRWNAI
		ŀ				MTVLRASKKDLELGGHMASFQ
						SSATIYDVCFNHFFRARNEQDG
		İ				GDLVYFQGHISPGVYARAFLEG
						RLTQEQLDNFRQEVHGNGLSS
						YPHPKLMPEFWQFPTVSMGLG
						PIGAIYQAKFLKYLEHRGLKDT
						SKQTVYAFLGDGEMDEPESKG
		İ				AITIATREKLD/NLVFVINCNLQ
İ		•				RLDSP\RWDELLRKDTSGKLIQL
						MNETVDGDYQTFK
17768	48136	Α	17874	452	1395	LAGATGTAIRLILHAPLLLPIKM
						RLP/TVSEGKPDSIPAAEKFAAE
					ļ	NKNTYGALASLELAQQFVDKN
						ELEKAAAQLQQGLADTSDENL
						KAVINLRLARVQVQLKQADAA
						LKTLDTIKGEGWAAIVADLRGE
						ALLSKGDKQGKHQRAENCEIQ
						GYIDMIKHLVAPLVFTSLILTGC
						QSPQGKFTPEQVAAMQSYGFT
						ESAGDWSLGLSDAILFAKNDY
						KLLPESQQQIQTMAAKLASTGL
						THARMDGHTDNYGEDSYNEGL
						SLKRANVVADAWAMGGQIPRS
						NLTTQGLGKKYPIASNKTAQGR
						AENRRVAVVITTP
17769	48137	В	17875	1	1579	· ·
1 7770	48138	Α	17876	182	2069	

SEQ ID NO:			SEQ ID NO:	Nucleotide location of first		Amino acid sequence (X=Unknown,
NO:	of peptide sequence	noa	in USSN 09/540,217	codon for peptide	of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		02/340,217	sequence	or peptide sequence	deretion, a possible nucleotide insertion,
17771	48139	Α	17877	I	2017	MESGVATRPIADFDVYIDKLTE
						FVYKTNLFMKPIFSQARKAPKR
						VVLPEGEEARVLHATQELVTLG
						LAKPILIGRPNVIEMRIQKLGLQI
						KAGVDFEIVNNESDPRFKEYWT
						EYFQIMKRRGVTQEQAQRALIS
						NPTVIGAIMVQRGEADAMICGT
						VGDYHEHFSVVKNVFGYRDGV
						HTAGAMNALLLPSGNTFIADTY
						VNDEPDAEELAEITLMAAETVR
						RFGIEPRVALLSHSNFGSSDCPS
						SSKMRQALELVRERAPELMIDG
						EMHGDAALVEAIRNDRMPDSS
						LKGSANILVMPNMEAARISYNL
						LRVSSSEGVTVGPVLMGVAKP
						VHVLTPIASVRRIVNMVALAVL
						FVNADETTVVNFHACFACVEV
		l	;			FTVRHTTNRYQHGVVTLRFSG
						CFFAFHRHINAVFFRFNIQAVFV
				ļ		ALRPEVIAIMHKLREQGHRVVV
			i			LSNTNRLHTTFWPEEYPEIRDA
		•				ADHIYLSQDLGMRKPEARIYQH
						VLQAEGFSPSDTVFFDDNADNI
						EGANQLGITSILVKDKTTIPDYF
				,		AKDKARHRTRPLWAWLKLLW
						QRIDEDNMTTLAGNLAYVSLLS
						LVPLVAVVFALFAAFPMFSDVS
						IQLRHFIFANFLPATGDVIQRYIE
		l				QFVANSNKMTAVGA\ASLAISS
					1	YLLSLIANVLMLMS*SVLRMS WNPKWAWSSIWYS
17772	49140	<u> </u>	17070	31	212	RGTGHCALYRRAAARCAKITL
1///2	48140	Α	17878	31	312	
						TTFISLFSSYFVIYNILLLIYLIFA
	·					ASFCLLVV*CIITVFITVCSAGD
						VSGMSATWCAFVAISVDHFTG
1	10141	<u> </u>	15050	•	1000	FNRF
17773	48141	A	17879	1	1098	
17774	48142	A	17880	1	1015	
17775	48143	В	17881	1	1917	
17776	48144	A	17882	931	1773	
17777	48145	Α	17883	785	956	KKWMHSAV*WRKRSIRRVSSL
						GY*TQAKDRRFALPELRRIVCS
						TVRRYVRRWRTNRT
17778	48146	A	17884	221	367	
17779	48147	В	17885	1	1318	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
17780	48148	A	17886	1190	1634	APSQTVSLPEVASSMSGVMAG
		-				VEAVSSPGPYAVCAGSLLPSGW
						LCVKTSVIAMIASAWSAPGAVS
						IKRSSAVSVRIWALCGA\QPECA
						VRSLPSYPAFRQVLAPVTHAL*
						PRRSSGCP*CDYSLQLLLQ*PAH
						*HLQGLLRKQLSCQR
17781	48149	Α	17887	1	2499	TIEQUEEKKQESCQK
17782	48150	A	17888	454	1698	
17783	48151	A	17889	635	1665	KSIFWRAPADCEP/QVRQSAKV
		1	1,,003			QFGDYQANGMMAVAKKLGM
					ļ	APRQLAEQVLTHLDLNGIASKV
						EIAGPGFINIFLDPAFLAEHVQQ
						ALASDRLGVATPEKQTIVVDYS
	ļ.					APNRGFFGGGAPLGTTLQDLTR
						PTKEDGLQAKPISFWASFALQF
ŀ		1				VNVKIILYGVTALSTFVLPQTQ
						ALSWVVGVSVLLAMIGTFGNV
						CWALAGHLFQRLFRQYGRQLN
İ		1				IVLALLLVYCAAFSKGTSDNVA
]				LLFRSQRVKANCITGYTDGQLR
İ		l	ļ			
	}	1				IFFRMLNCIFQSFATQNVNVQV
	İ	1				LTTFNANFWFHFYWNFTVFNV
		1				TQFAYCNHFIFCITCFCNDATPF
17704	40150		17000	72	212	AFFAVQQPERVHRKTAILL
17784	48152	A	17890	73	212	VEDVEL DIVVEWUNDCAMATA
17785	48153	Α	17891	2049	2445	VFPVFLRIYVFWKKPGMKVTN
		1				ARNWQLKPLFTELPVILPDTQL
						HYVAWMELYSPAE*ERIQA*FV
	}]				VWSWNIWLY*A*RLIQK*IIALT
]				PVVRTGKTKLMSAILFNITIHR
17706	40154	_	17002	1	1476	MKAMNLSSPKRRLPPRNCGKK
17786 17787	48154 48155	A	17892 17893	332	1476 572	RGGA*RQKHRRTSALSLLSLP*
1//8/	48133	A	1 / 693	332	372	NMATAVHLHRFSTRRAVTSSS
		1				
						ARKWTNSGAMSGLNRASA\AV
17700	10156	_	17004	100	405	LRV*QSPEDGCCARIR PRNLRKSGPPISDAISPTSKP*TI
17788	48156	Α	17894	80	403	
						TEPISAPSLLATYKAAGCGGTT
						QCTAINAVHSGMASFNNEVLV
						FLAIEKARGINSTTPTSTNSVMP
17700	40166	<u> </u>	17005	 	1202	QIRPTSTIMTSTESQRQR
17789	48157	A	17895	1	1392	
17790	48158	A	17896	1071	1374	
17791	48159	A	17897	1971	2306	
17792	48160	A	17898	209	449	
17793	48161	A	17899	1	240	
17794	48162	A	17900	1	456	
17795	48163	В	17901	222	3807	
17796 17797	48164	A	17902	3140	3219	
	48165	A	17903	240	298	ı

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
17798	48166	A	17904	1902	2047	NQPVYRSNGD/RIWRTGKLFPT TVFRLIRTLSPRAFPPFTSYRRCS VAY
17799	48167	В	17905	1	1994	
17800	48168	Α	17906	1	2151	
17801	48169	A	17907	3	1127	GTPYASQPVIRPVEGSKPLIKQV LDIGAHGTDKTIAQKLARVAR MHFARQRLAAVATFACSAQNA IALMEEIAANFSYEMIRLTECIL GFTWNRLYQGINVHNAERVRQ LAHDGHELVYVPCHRSHMDYL LLSYVLYHQGLVPPHIAAGINL NFWPAGPIFRRLGAFFIRRTFKG NKLYSTVFREYLGELFSRGYSV EYFVEGGRSR/TGRLLDPKTG\M LRGGTRPITLIPIYIGYEHVMEV GTYAKELRGATKEKESLPQML RGLSKLRNLGQGYVNFGEPMP LMTYLNQHVPDWRESIDPIEAV RPAWLTPTVNNIAADLMVRIN NAGAANAMNLCCTALLASRQR SLTREQLTEQLNCYLDLMRNVP
17802	48170	A	17908	192	486	PVLVYLWVSFSWLCGRSLVCD AFGCVLCVLCFWCGVCWGWF SRCLVLVRVLDCVAVAWVCPL GVGIGLWV/RVLPTGLVSAVPC CCDVLARFRVDLCWG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ŀ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
17803	48171	Α	17909	1	2470	MFQPSDSGKSFIFNMSVGYNLE
		ł				GIKQPPMQQFIDNMMDASDHP
		ļ				KFAQYRDTLNKLLQDDAFLAR
		1				HGLQEKRESLQALPARIPTSMV
		1				HGVTLSTMHGCPPHEIEAICRY
		1				MLEEKGLNTFVKLNPTLLGYA
		1				RVREILDVCGFGYIGLKEESFD
		ł				HDLKLTQALEMLERLMALAKE
		ł				KSLGFGVKLTNTLGTINNKGAL
		ļ				PGDSTSGVLRTYERLLTAWML
		ļ				TCPCADEHQRGESVKLATALT
		İ				WWLRGSQNRDEITKDGFQSNH
		1				AGGILGGISSGQQIIAHMALKPT
						SSIDVPGRTINRFGEEVEMITKG
						RHDPCVGIRAVPIAEAMLAIVL
						MDHLLRQRAQNADVKTDIPRC
ŀ			}			AQSIGSFSNGCIVGADTLPIQSE
						HYQVMRTDQRRYFGHPDLVM
						FIQRLSSQVSNLGMGTVLIGDM
						GMPAGGRFNGGHASHQTGLDV
						DIFLQLPKTRWTSAQLLRPQAL
						DLVSRDGKHVVSTLWKPEIFSL
						IKLAAQDKDVTRIFVNPAIKQQ
						LCLDAGTIATGCAKCDPGVSIG
						AYAYLLTLRAQCLAHDLPDPLE
İ						PLEIDGTLLPRYVFIHGGPRVFT
		İ				YYTPKEESIKLFHDYLDLHRSN
						PNLDVQMVPVSVMFGRAPGRE
						KGEVNPPLRMLNGVQKFFAVL
		Ì				WLGRDSFVRFSPSVSLRRMADE
						HGTDKTIAQKL\ARVARM\HFA
]		ŀ				RQRLAAVGPRLPARQDLFNKL
17804	48172	Α	17910	1	734	
17805	48173	Α	17911	1	962	
17806	48174	Α	17912	1630	2633	
17807	48175	Α	17913	1360	1600	HSTGRWATTINVPITCGTKARH
			<u> </u>	1		STSSICKTGMILLWANALKQHC
				1		LLSRNLP*KGESINVANSLLSYF
						NLRPQCNQTRTQD
17808	48176	Α	17914	1	816	
17809	48177	A	17915	703	1546	O LUD PROTECTION OF THE PARTY O
17810	48178	Α	17916	209	485	QVIRPTFIPSTSSNGAISLPASTR
						RDS*MANAS*RCSLA*RATMFP
]				A*SATSIFHARPGVEGT*IVRIV
			<u> </u>			VIMLRLLSRRKVRGIRLPAAVLI
17811	48179	Α	17917	619	1167	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
17812	48180	Α	17918	1	764	DSGQQLDDVQLARYLLSGQNQ
			•			PKSDRPYRVR*LTGAMLKLILT
						L/VIEGNVTLGHRVKIGTGCVIK
						NSGDWR*LRNQSVYRCGRCES
						GSGLYHWARLPVCVLVLSCWK
						VLTSSLVPPD
17813	48181	Α	17919	3	1462	
17814	48182	Α	17920	122	983	
17815	48183	В	17921	1	1287	
17816	48184	A	17922	289	958	
17817	48185	Α	17923	1	5079	
17818	48186	A	17924	1 802	255	TECD CADC A COUNTY CTERVISE
17819	48187	A	17925	1892	2448	TESRCADGAGSVDVGTEPVVP
						RVELVAKWPGSRAGGTGLGCS
						PWVGAGCVGGLGPQARLSGPT
		[ASTEASAKAASTLASLMWLRV RL*GACPFARTGATAPSSSKGP
						ELGVSGLRKATWMGRGSGCCT RGLGAGEWRAVPESCGLDESS
						EDSDDRDHAVPFSSSSCRRSML
						RPRRVARGLCRRRGV
17820	48188	В	17926	52	888	RPRRVARGLERRRGV
17821	48189	A	17920	929	1148	TVWNGTCGLPFSSLMKCMVIM
17621	70107		11/2/	122	1140	IMRAIQKKMMSKPDTITLVGW
						N*RRASVFSGQPRVEKVHSAEE
						NQVSRTSSS
17822	48190	A	17928	125	937	RISMVFPWCFCSCSRTLATTPAP
		ļ				TVRPPSRIAKRRPSSIAIGWIRVT
						TILMLSPGITISTPSGSSMVPVTS
						VVRK*NCGR*PLKNGV*RPPSS
						LDRMYTSDSNLHGLRPAVSPVT
						NTPA/SDGYYSHDGSLAQVDLS
		İ				ANYHEGQYTSAGLSLQGGATL
		ŀ				TTHGGALHRTQNMGGTRLLID
		ŀ				ADGVADVPVEGNGAAVYTNM
						FGKAVVSDVNNYYRNQAYIDL
						NKLPENAEATQSVVQATLTEG
						AIGYRKFAVISGQKAMAVVNF
		İ				AEGKVSLAVV
17823	48191	Α	17929	3992	4213	VPAIQQKFPRSWKLGGERPPKA
						FRLRLSAA*RKSVP*AQRSLAD/
						SGMPLRLEKCLDTF*TQIV*SAF
						KIQREGS
17824	48192	Α	17930	1	1636	
17825	48193	Α	17931	1	900	
17826	48194	Α	17932	1	939	
17827	48195	A_	17933	427	1250	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
17828	48196	A	17934	521	805	LCPSNTSPLSCLLCDLVASTEM APMRVFQPAPKVNA**LFATIEP PYSVNKLLKPY**PAWTKSLCL LAPSIWLRPAAKSYFDTPGCSIF GTAA
17829	48197	Α	17935	1	2052	
17830	48198	В	17936	1	573	
17831	48199	Α	17937	1	1278	
17832	48200	Α	17938	161	659	
17833	48201	Α	17939	1	588	
17834	48202	Α	17940	695	1281	KLKPYSWAFSRQSGRSVILTPPR VTVVVVAGFGPIPVFGTVMVRP GWAAP*FLPVASVYFP*AGLAG FLKKNRGCCANPAPPPIRRPIMY PP/MQRLQEPPADWWACAWFS GKD*\WKKSPPNCTRCA
17835	48203	Α	17941	261	2045	
17836	48204	A	17942	1	637	
17837	48205	Α	17943	31	280	
17838	48206	Α	17944	1	582	
17839	48207	Α	17945	1	2496	
17840	48208	Α	17946	330	507	
17841	48209	Α	17947	1087	1875	
17842	48210	Α	17948	1	476	
17843	48211	Α	17949	1	537	
17844	48212	A	17950	2	492	
17845	48213	Α	17951	1	1101	
17846	48214	Α	17952	2538	2833	TRRNWRPIQSSEESWRAAGSST PLTTQNRQQTAGIY*RSR*TAA RR*SPALPRRNPRLATAG\WQY ATQTRQQ\RVPQL*LSAPATRK NRLLANAWLTM
17847	48215	Α	17953	125	1831	
17848	48216	A	17954	2	825	
17849	48217	A	17955	1	840	
17850	48218	Α	17956	1	1725	
17851	48219	A	17957	463	1824	
17852	48220	A	17958	2	117	
17853	48221	A	17959	1	2640	
17854	48222	В	17960	223	1560	
17855	48223	A	17961	638	746	RIP*MT*WS*KPWHKNYMMHA QASVDNLIKWKKGYQ
17856	48224	Α	17962	385	514	
17857	48225	A	17963	17	225	SSPSKFTVAFFASLVMASAVSL ALSLRSLTTLSI/CGLVLQEERT DDALVEQFSPIRRARGHAPQQE ATEM

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
17858	48226	Α	17964	1	521	LVFLVLFVGFLSCFKSTYTLTY
		ļ				GDSPSLCDGLLII*IISRIDTNKPA
						DNSISGIRDVGWVAKSYQVA/V
						GERLHNRTPGTSHQSDTRFDF*
1						LC*PGSACPNLVSEHERPQANM
						LVEASGRLAA*YQNALSCSFGY
						GVGLVRRQDFSSLGRFLQSAFA
					·	SNRY*QVVWVFEFQQVS
17859	48227	Α	17965	267	554	LRFRFTRAVGRIFHRLDLLVLL
						GCNRYGRRGGDHGLCSVLVPR
			1			SLRLGRLPGGDSAAADAQSRH
i		ŀ				REKCR*DGVLVCDDQNRQTPV
		ŀ				WKPHQEDVQV
17860	48228	Α	17966	1	915	
17861	48229	Α	17967	1226	1399	SDSNISICTCAQIH*KHFRII*NCT
		į				L*EVEP*RLS*M*LQRKCVQGEP
! !						SKVHSLWGI
17862	48230	Α	17968	828	1070	SFFDYQTSALEHARR*LHCRT*
						NAGWLPHRTDRSSGYHGRTCS
			}			AAGRCSSGDHSSYQQQNPAEK
ł						LPAGHPQRSGSPERTG
17863	48231	Α	17969	45	1198	
17864	48232	Α	17970	348	508	RLFRWCPM*S\WASYFPPESQSL
						PDSYWLF*ECNLPTAYNSQNAII
						RESHAIYS
17865	48233	Α	17971	251	601	MAKKMSLSIAVFWRRLYAY\Y
						PHLR*KS*TNW/QWWRSALS*A
1						V*PNRS/IDSRWSIIWSVLFGML
						V/AT/ALAPAFWLRY*KKSFSA
1						WNPTKSPRCLSNARPCCSLSKK
						ASLPWTIAARSR
17866	48234	Α	17972	1	2121	
17867	48235	С	17973	1	1710	
17868	48236	Α	17974	1	1335	
17869	48237	Α	17975	1	1497	
17870	48238	Α	17976	2	824	
17871	48239	Α	17977	I	1965	
17872	48240	Α	17978	1	3555	
17873	48241	В	17979	197	2021	
17874	48242	Α	17980	1	348	
17875	48243	Α	17981	1	891	
17876	48244	Α	17982	1	679	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
17877	48245	A	17983	3	2502	EAIREVSNPI\PQEEIWTHREAVP
						PVQLLTGWCVTAKAPPDISAIS
		ļ				ALTAVKHGNCSSLTPLLNPPGS
		ļ				DVIVCAEMDEQWGYVGAKSR
						QRWLFYAYDSLRKTVVAHVFG
						ERTMATLGRLMSLLSPFDVVIW
						MTDGWPLYESRLKGKLHAILV
						TFEEKFQEEPVKKQIIMVSISVS
						DTGGKKEKPKLINLSSKTNGLQ
						NTTEAQLCETKSDFELSFSGFNP
						ISIDTEDMKTGCCFQEQSVRRA
		İ				YFKATSSPRRGLIITKLTTRVDG
						YVADDPSRRYSGHGDLLWPNA
		ļ				GNSCVETTVGSLLCEVQAAPTP
						`
		ŀ				HTLEGKHGCRNKEKKRLTHSG
1			1			ACTYTFSELGLELWAWNCGKT
						EVNLGAVGKGVGGDRGFKKIH
						AKSQFTCSVLYTFTIKQGISLTL
						WSKVMHILEHLKENSKQLTSIC
						TRYQRVFGPTWLHETIFCKVAL
						TATLLAELEKTSTLQTFLDKIWI
						LGAQQHTGSQEPSQALLTCDPA
						WKKALRKVTQKGGAFFGVDN
						LGSSLSPTTDLLYEEQRLTSLSG
						SWFPICVMVELTLRAGSSPGGR
						LYVEFDQAVTKEPRFLQYKHY
						WDGAVAGSSILRQDSFVAEAST
						TITRIESPSVPSRSSTFLDDFSITQ
						QHRRHAGLSVMLSLIISPPVRGL
						CQERTLRRDFRERLMYLDLEY
						QSGYCRLRTAGSEMQMVPDVL
						IFAHRSGLAGLFPGLDRRLVNA
17878	48246	Α	17984	10	211	GSQSTIFTFAARW\NLWANENT
						DTSISPLGVGEQRHCSSP*RFPD
						SDRCSPWSPRSQGQNQSPQTW
					ł	Q
17879	48247	A	17985	1	1725	
17880	48248	A	17986	1	379	
17881	48249	A	17987	3	273	
17882	48250	Α	17988	3	1157	
17883	48251	Α	17989	76	378	
17884	48252	Α	17990	1527	2005	
17885	48253	Α	17991	578	1312	
17886	48254	Α	17992	1	1758	
		Α	17993	1	2433	
17886 17887	48254	-		ł. ———		

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
17888	48256	A	17994	. 1	537	SQEELLQQALVAQLA/HSEQLE GVQLPEPDEPSARHALPANEPRI VLNNGVVSYNDRPILNNLSWQ HIGYVSSSLHLDYRVSTTVRNV ILSGYFDSIGIYQAVSDRQQKLV QQWLDILGIDKRTADAPFHSLS WGQQRLALIVRALVKHPTLLIL DEPLQGLDPLNRQLIRRFVDVLI
17889	48257	Α	17995	1	536	
17890	48258	Α	17996	1	1319	
17891	48259	С	17997	74	341	
17892	48260	Α	17998	3	103	
17893	48261	Α	17999	1	745	
17894	48262	Α	18000	1	663	
17895	48263	Α	18001	2	297	
17896	48264	A	18002	8	340	VPANSLSAIALFRRESVTPLAAL AASPVCAGRFVLLKNHRRDLP* ISVPVNSSSNSARSLVWHSGRQ QIAPDWSSKDDFMRLGKPNRFS VLLLATWIR*STVIRRSPFTS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
17897	48265	Α	18003	1	1341	MVHRFGIRVRFVALGFYQDVIL
						PVRAHRHEWGHYVHLCASIQR
						QIDVSFARDATSHSSCPVTGVR
						FWGYPFAANKVVIAGFELEGET
						MTRPGLICAEGGLLVMIVSLRS
				ŀ		ELYDSYHEFTRQRVTEIVEKAA
						NHLPAVVEMMTLHRCAINRHN
						FVRRVGPAGISTFFASSPDGVN
						AGFSTFRVVFTCQRSRPLQRHS
						HADVQTATLLLNTVNLEQKQA
						NAILSGLSDMIPNSSPESAPEIQL
						LQSRMILGKTIAELNLRDIVEQK
						YFPIVGRGWARLTKEKPGELAI
						SWMHIPQLNGQDQQLTLTVGE
						NGHYTLEGEEFTVNGMVGQRL
						EKDGVALTIADIKAKPGTQFVL
						SQRTELEAINALQETFTVSERSK
						ESGMLELTMTGDDPQLITRILN
						SIANNYLQQNIARQAAQDSQSL
						EFLQRQLP*TLSKNRPTPFSAA*/
						DDMIPNSSPESAPEIQLLQSRMI
						LGKTIAELNLRDIVEQKYFPI/D
						GSRLGEINQRKTR*AGDQLDAY
						STTEWSGSATDT\QVGENGHYT
						LEGEEFTVNG/NGRPASGKRWR
						CADYRGH*GQDQQLTLTVGEN
						GHYTLEGEEFTVNGMVGQRLE
						KDGVALTIADIKAKPGTQFVLS
						QRTELEAINALQETFTVSERSKE
						SGMLELTMTGDDPQLITRILNSI
						ANNYLQQNIARQAAQDSQSLE
						FLQRQLPEVRSELDQRKKNSTFI
17898	48266	Α	18004	414	893	VFAQHGLTVAAIAAHARDRWF
						VGRTAHANPPVRCSTDGSGRVP
						VTRYSGFVAAAVILLTASPRGG
						ECSREAPLCSVSEYLLLQSAGE
						YAPAPRCVLRCVPSVDEDFVPV
						VPD*APPVEVALFLLRLTPVAR
Í						PADGGWLLAAIRPDALTSLAIV
: }						VLAAAR
17899	48267	A	18005	432	1108	
17900	48268	В	18006	1	753	
17901	48269	Α	18007	46	282	
17902	48270	A	18008	1	1201	
17903	48271	A	18009	1	618	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
17904	48272	A	18010	1	1045	MLCTKWDKLGLEKNAGKNKN AYKWYGWPANAPDEPHIEQLV GECRVMRQLKRLISRIAPSPSSV MVVGESGTAIPVVSHAEFKGGF ADIGVHYLDWTSRTTEKSSTKS HKDDFGYLEFEGGANFSWGEM YGFFDWENFYNGRHNKPGSEQ RYTFKNTNRIYLGDTGFNLYLH AYGTYGSANRVNFHDDM\FLA GFGFQF*PAGGW/WGSNPFFAK\ RYTRSKPYYT\GDNGYV\AGWV AGYNFMLGSEKFTLTNWNEYE FDRDATYAADFLPLYDVDCQD NGNLEYDTYSQPEWKHNLFDH YLAVLYRFKDESGKEQFSGAV VKTREATPGKEIEAITRRMLDFS PRLKKLA
17905	48273	Α	18011	228	430	YPRQHRQRADGSDRHR**/CWS A*IWANPTSNLPPCRFALTKRK RPILCAPPSRQSYAAWCRWEIRI A
17906	48274	A	18012	296	1178	
17907	48275	A	18013		1395	MPSDISLASCRLRATSRLFTDHR RQSDCMTAINRIPIVDDEDNVR RFETHCANNVRTALHLFADIHP DVVLMDIRMPEMDGIKALKEM RSHETRTPVILMTAYAEVETAV EALRCGAFDYVIKPFDLDELNLI VQRALQLQSMKKEIRHLHQAL STSWQWGHILTNSPAMMDICK DTAKIALSQASVLISGESGTGKE LIARAIHYNSRRAKGAFIKVNC AALRESLLESELFGHEKGLFER ANEGTLLLDEIGRM/PLVLQAIL RILQEREFERIG/GHQTIK/VDIAS LLAPTRLQAMVKEGTFREDLFY RLNVIHLILPPLRDRREDISLLA NHFLQKFSSENQRDIIDIDPMA MSLLTAWSWPGNIRELSNVIER AVVMNSGPIIFSEDLPPQIRQPV CNAGEVKTAPVGERNLKEEIKR VEKRIIMEVLEQQEGNRTRTAL MLGISRRALMYKLQEYGIDPAD
17908	48276	Α	18014	1	771	
17909	48277	A	18015	1	2154	INTOFTH DOLODGED AND HOLES
17910	48278	Α	18016	859	1014	IKTGETHPGIGPSVM*RHSKAD LTHRHCGGIPEPQSVSFGSSQH GGGQFPPL

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
17911	48279	A	18017	1224	1520	SPLVNQ*GA*KFFSRGNLFPQG ETH*KTKGVPCWTRYPDLKRS DSAEIFKTPANASGRTSGSLGC AAQLEKVADGRDLDEKLAEAG IGGSNKTSVWQG
17912	48280	В	18018	182	2392	
17913	48281	Α	18019	72	266	
17914	48282	Α	18020	600	914	PYRGSPRNGHQLCDHCSHRLSE DPFVRYSNHCQSISNYRCDKPP VRPLSPLHQKTDRKHHQRPDRP FQWSAAPAHGRWRSGDS*ETP LHYAQPAGKLRWLWVNY
17915	48283	A	18021	1102	1171	LLGYGQQPPASNLLSEG*RSVV L*ISALRM*ALPPQKSSDAPNG WLPNRWPISSTHMIR
17916	48284	Α	18022	1	1279	
17917	48285	Α	18023	1329	1403	
17918	48286	Α	18024	442	918	
17919	48287	Α	18025	3	1345	
17920	48288	В	18026	436	988	
17921	48289	Α	18027	637	1086	
17922	48290	A	18028	1	321	
17923	48291	Α	18029	1	845	
17924	48292	Α	18030	298	581	GARAEARAGSGQGAGLGVSSP VRWRGWSDKGAERPGGPLSPL REVSRAGPSG/HAARGQQPGRP RFPPPGPRPPRRRCPCTRARPGG AAGGGPCL
17925	48293	С	18031	450	604	
17926	48294	Α	18032	1279	2253	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
17927	48295	Α	18033	1	1957	MAQFHVSHPPAEWPRHVLLLM
						ADSKNELQNMNTGQTCGTRQT
						LKSKLQGWNKPGIQSDPGSQG
						HPRCLKTFPLVAQSSTQERNHV
						MEKAVLKSONWAIIEEFITKIG
						WFPDYSGKVKYLSSPKSSCTTV
						LQVTHFSLFFHPNPDELVSKIKV
						WSKHRLYQNNSEAFLEVQIPEP
						KCEVYSMRTMGRRQPGTAMD
						LNAGGSICNVPRTLLHLTGESTF
				1		RDRQRVGAYLDIRGWYILVLV
						VDKEKLGIPGQKQNSSKILFGSE
			j			KLKQWVYVKNHAKESITQEAM
						GIWVGQQRKRQPLGYEERKLT
						NRKDIHIKNPSVCHHHQRPKVD
						KTTKMGKKQSRKTGNSKNQST
						SPTLKESSSSPATEQSWTENDFD
						ELREEGFRRSDYSELQEEVQNN
						GKEVKNFGKKLDEWITRITNAK
						KSLKDLMELKTKARELCDERTS
1						LSSQCNQLEERVSVMEDEMNEI
						KQEEKFREKRIKRNEQSLQEIW
1						DYMKRPNLRLIGVPESDGENGT
-						KLENTLQDIIQENFPNLARQANI
1						QIQEIQRTPQRYSLRRATPRHIIV
						RFTKVEMKEKMLRVAREKGRV
		1				THKGKPVRLTADLLAETLQAR
						RQDTHRLKIK\DGGRFTKQMEN
						RKRQGLQS*SQIKQNLNQQRSK
						ETKKAIT*W*RDQFNKKS
17928	48296	Α	18034	1	2838	
17929	48297	A	18035	1	897	
17930	48298	Α	18036	26	308	
17931	48299	Α	18037	749	1020	
17932	48300	A	18038	1	1773	
17933	48301	A	18039	1	1890	
17934	48302	Α	18040	1	756	ODDE AVA (DDD LEDVINE) (TODE
17935	48303	Α	18041	399	660	CPPEAVMPRRHEPVWEMTGFF
				1		LPDVLLSDSSDTSEGFRLHLET
						WAPFLCRCLSLRLIS*HHHQTT
17936	48304	A	18042	909	2005	GNSSPPVSASQHQSVDRDSLN
17937	48305	A	18042	1	490	
17938	48306	A	18044	104	775	
17939	48307	A	18045	170	451	
17940	48308	A	18046	1	207	
1,770	1.0550	l* *	1.00.10	I		

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
17941	48309	A	18047	1146	678	LGFLLRLSEMPRKQGDYRTRIW
1.,,,,,	1.030)	[10017			KFEDGLSNVLVIQLNKLIICVM
						CLVRDCDVLKTYFHR*SGLLPI
1	1					KDTS/RPFTRVTVQWGKKNDQ
						TFQGLLDTGSELTLIPEDPKRHC
	1					GPPVKVGAYGGQVLHSKWAPI
	l					YHFLSSHNCNGFCYISIANLPKK
						FGQNIKECLTNNRIVPFNCVQD
17942	48310	A	18048	270	1444	I OQNIKECE INNIG V FINC VQD
17942	48311	-	18049	700	1062	
1		A	1	<u> </u>	1096	
17944	48312	A	18050	3		
17945	48313	A	18051	1	1500	MEDAN ICINIMINAL AVODCEDE
17946	48314	Α	18052	1	6215	MEDNLISINKIKILLAVSDGEIDE
			}			TFSLKQLMFNSVPVQNEDGSFN
	İ					FEGVKAEFRPGTQTQEYIKGME
	1					DSSSEVTVNREVTTDNPYTISVT
		1				NKTLSAIRIKMFMPRGVRIESN
		ł				GDKNGVRVEYEVQQAVDGGSF
		į				ETVLTDVIEGKTMSGYDRSRRV
		1				NLPNFNNQVIFRVVRKTPDSND
		l				SNVVDAIQVRSYAEVIDAKFRY
						PLTGLLFVEFDSKMFPNQLPTIS
						IRKRWKIVNVPSNYDPESRTYN
		l				GNWDGTFKKA
17947	48315	Α	18053	1325	1417	
17948	48316	Α	18054	1	4971	
17949	48317	Α	18055	251	1091	
17950	48318	Α	18056	1	2277	
17951	48319	Α	18057	2	448	KNDKGDIIVTTKSGGRGTSTVS
						FKLLKPEKI/GSKFSQKDLEMLF
	1					HGMRADFTSENFSAAWYLIEN
						HSNTSFEQLKMAVTNLKRQAN
						KKSEGSLAYVKGGLSTFFEAQD
						ALSAIHQKLEADGTEKVEGSMT
						QKLENVLNRASNTADTVYE
17952	48320	A	18058	124	984	

SEQ ID			SEQ ID NO:		Į.	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		05/540,217	sequence	or pepular soquenes	, possible same same same same same same same sam
17953	48321	Α	18059	142	1393	DKKPGMVPPPPGEESQTVILPP
						GWQSYLSPQGRRYYVNTTTNE
						TTWERPSSSPGIPASPGSHRSSLP
						PTVNGYHASGTPAHPPETAHM
	İ					SGRKSTG\DSQNLGSSSPSKKQS
						KENTITINCVTFPHPDTMPEQQL
						LKPTEWSYCDYFWADKKDPQG
		l				NGTVAGFELLLQKQLKGKQMQ
						KEMSEFIRERIKIEEDYAKNLAK
						LSQNSLASQEEGSLGEAWAQV
						KKSLADEAEVHLKFSAKLHSEV
						EKPLMNFRENFKKDMKKCDHH
						IADLRKQLASRYASVEKARKAL
		İ				TERQROLEMKTQQLEIKLSNKT
						EEDIKKARRKSTQAGDDLMRC
		İ				VDLYNQAQSKWFEEMVTTTLE
		ł			1	LERLEVERVEMIRQHLCQYTQL
						RHETDMFNQSTVEPVDQLLRK
				ł		VDPAKDRELWVREHKTGNIRP
15054	40222	 . 	10060	,	425	VDMEI
17954	48322	A	18060	12	926	
17955 17956	48323 48324	A A	18061	3	693	RPPLAYLRDGDARAGLNGLQL
17930	46324		18002		093	AVLAMLSSRKMFCKKSGQSYS
						PSRVLITENDVKEGLQRSHILYD
		ŀ				RAGEEHYNCISALHKSMRGSD
		ŀ				QNASLYWLARMLEGGEDPLYV
						ARRLVRFA\SEDIGLADPSALTQ
				1		AVAAYQGCHFICMPECEELLA
						QCVVYFARAPKSIEVYSAYNN
		ŀ				VKACLRNHQGPLPPVPLHLRN
						APTRLMKDLGYGKGYKYNPM
						YSEPVDQEYLPEELRGVD
17957	48325	В	18063	1	2020	
17958	48326	A	18064	19	404	
17959	48327	Α	18065	1	407	FRATITSAKMGAYKYIQELWR
		1		1		KKQADVMRFLLRVRCWQYRQ
				1		LYALHRAPRPTRADKARRLGY
			1	1		NDKQGYVIYRIRVRRGGRKRPD
						PKGATYGKPVHHGDNQLKIAR
						SLQSDAEEPAGRHCGALREQSS
				1		Y*VGEHST
17960	48328	A	18066	53	431	AKKGAYNYIQELWRKKRADV
				1		MRFLLRVRCWQYRQLSALHRA
				1		PRPTRPDKARRLGYKAKQGYVI
	1	1		1		YKIRVRRGGRKRPVPKGATYG
				1		KPVHHGANQLKSAQSLQTGVQ
						ERPGHP*YAMRVLKAYRARPD
17961	48329	Α	18067	5	328	
		•	<u>. </u>		•	

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
17962	48330	A	18068	93	836	AKMGAY\KYIHELWRKKQSDV MRFLLRARCWQYPALHRAGTE WQLSALHRAPRPTRPDKARRL GYKQ\AKQGYVIYRIRVRRYI*D SCSPWWLKTAQFPKGCNFTGK PVHSWVFNQLKSFASKAFQFR WQEGGEAWDGTCGGLLEEFLN FFNWVWV/GKIST*QNFFEGLIL I/VIPFHKS*SEENPDTQWDQPN QSHKHR\EMRG\LTSARPEKSRG L/GKKGHKFHHTIGG/SLRRAT WRRNTLQAPPLPLI
17963	48331	A	18069	1495	2055	
17964	48332	В	18070	125	2530	
17965	48333	A	18071	345	800	RWRLQNLSFCSSCSGSGSEGAT CFARGR*RPPGNSLNVSRMGTR SSYSCRRHSPAFLSWGKGRCHL CNKAHLLAGA*HYSFLSGSAPY *N*VNGQI**R*TAHRSTPDAPT GH*NCCS*L*CHSY*QRA*PGY RHD*CRMCC*CADCSHAC
17966	48334	Α	18072	181	1687	
17967	48335	Α	18074	205	1734	
17968	48336	Α	18075	1	705	
17969	48337	Α	18076	590	685	
17970	48338	В	18077	1	881	
17971	48339	A	18078	1043	1248	ITLGIMAIFTVLILPIHEHGMVFH LFVSSFISLSSGL*FSLKRSFTSL VSWIPKYFILFEAVVNGSSL
17972	48340	В	18079	1	1977	
17973	48341	A	18080	805	1153	AWKVLPFFVSFCF\LSSGL*FSL KRSFTSLVSWIPRYFILFEAIVN GSSLMIWLSVCLLLVYKNACDF CTLILYPETLLKLLISLRRFELLF LSMFSASFRSSCKAGLVVAESL
17974	48342	A	18081	471	1442	IWSFHIVPYFLEALFV*GVRKGS SFSFLHMAGQFSQHHLLSRESF PHCLFFSGLSKTR
17975	48343	Α	18082	188	355	
17976	48344	В	18083	1	1677	
17977	48345	С	18084	35	1816	
17978	48346	Α	18085	557	724	
17979	48347	Α	18086	179	346	

NO:	EQ ID SI	EO ID NO: I	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
17980	- 1					F .	*=Stop codon, /=possible nucleotide
17980	se	equence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
STPQQM*KNRNYN NQTRTQDEETHSK: QPAPE*ILGT*QPAPE*ILG					sequence		
STPQQM*KNRNYN NQTRTQDEETHSK: QPAPE*ILGT*QPAPE*ILG							
NQTRTQDEETHSKS QPAPE*LLGT*QNE NQREGNAHINES NQREGNAHINES NQREGNAHINES NQREGNAHINES NQREGNAHINES NQREGNAHINES NQREGNAHINES NQREGNAHINES NQREGNAHINES NQREGNAHINES NQREGNAHINES NQREGNAHINES NQREGNAMINE NAKENILAN NAKE	7980 48	8348	Α	18087	135	1461	INRIYILLSTTSHLFQN*PHSWK*
				•			STPQQM*KNRNYNKLSLRPQC
NQREQRHNIPESLG				1			NQTRTQDEETHSKSLNYMETE
**RRKERRIK*MQ* AKGVITTDPTEIQTT YANKLENLEDITOK LNQEEVESLNRPITE LPTKKRPGPDGFTA EGRIKYLGIQLTRD YKPLLNEIKEDTNK IGRINIMKMAILPKY KLPRTFFTELEKITI, ARIAKTILSKKNIAG LYYKATVYKTAWY QWNRTEASEITPHI EKNKKWGKDSLFN LAICRKLKLDFFLIP IKDLNVRPKTIKSLI 17981			i				QPAPE*LLGT*QNEGRNKDVL*
AKGVITTDPTEIQT:				•			NQREQRHNIPESLGHI*SNC*QD
YANKLENLEDTÜK LNQEEVESLINRPITT							**RRKERRIK*MQ*KC*RCNKN
LNQEEVESLNRPITO				1			AKGVITTDPTEIQTTIREYHKHL
LNQEEVESLNRPITO							YANKLENLEDTDKFLDTYTLPR
LPTKKRPGPDGTTA			i				LNQEEVESLNRPITGSEIEAIINS
EQRIKYLGIQLTRD							LPTKKRPGPDGFTAEFYQRYKE
VKPLLNEIKEDTNK IGRNIMKMAILPKK KLPRTFFTELEKIT ARIAKTILSKKNIAG LYYKATVTKTAWY QWNRTEASEITPHI EKNKKWGKDSLEI EKNKKWGKDSLEI EKNKKWGKDSLEI EKNKKWGKDSLEI EKNKKWGKDSLEI EKNKKWGKDSLEI EKNKKWGKDSLEI EKNKKWGKDSLEI EKNKKWGKDSLEI EKNKKWGKDSLEI EKNKKWGKDSLEI EKNKKWGKDSLEI EKNKKWGKDSLEI EKDLINVPRKTIKSLEI EKNKKWGKDSLEI EKDLINVPRKTIKSLEI EKNKKWGKDSLEI EKDLINVPRKTIKSLEI EKNKKWGKDSLEI EKDLINVPRKTIKSLEI EKNKKWGKDSLEI EKDLINVPRKTIKSLEI EKNKKWGKDSLEI EKDLINVPRKTIKSLEI EKNKKWGKDSLEI EKDLINVPRKTIKSLEI EKNKKWGKDSLEI EKDLINVPRKTIKSLEI EKNKKWGKDSLEI EKNKKW			i				EQRIKYLGIQLTRDVKDLFKEN
IGRINIMKMAILPK\ KLPRTFFTELEKITL ARIAKTILSKKIAG LYYKATVTKTAW\ QWNRTEASEITPHI EKNKKWGKDSLFN LAICRKLKLDPFLIP IKDLNVRPKTIKSLF EKNKKWGKDSLFN LAICRKLKLDPFLIP IKDLNVRPKTIKSLF EKNKKWGKDSLFN LAICRKLKLDPFLIP IKDLNVRPKTIKSLF EXTENSION A 18089 659 826 17983 48351 A 18090 3531 3878 17984 48352 C 18091 1 2793 17985 48353 A 18092 1607 1774 17986 48354 B 18093 1 1666 17987 48355 A 18094 1 1068 17987 48355 A 18094 1 1068 17988 48356 B 18095 1 1549 17989 48357 A 18096 557 623 KKWINYWTHTPSQ P 17990 48358 B 18097 1 1231 17991 48359 A 18098 263 1461 GGSSCYPSWEATIR KRILKVLTFLVLFF SHTMVATTWFFKC LKRLIKHRPCTOTTOP LEDDTYRWLSLE GFCLQRPFGDKQGT TGRDLTQQRQLRI DTIKELFRVYPGNV VGCRRCAVVGNSG PEIDSHDFVLRMNK DVGTKTTHHLVYP VSMILVPFKTIDLEX TISHTYIPVPAKIRV PAFIKYVFDNWLQC GILSVIFSMHVCDEX CILSVIFS CILSVI							YKPLLNEIKEDTNKWKNIPCSW
REPRIFFELEKITL ARIAKTILSKKNIAG LYYKATVTKTAWY QWNRTEASEITPHI EKNK WGKDSLFN LAICRKLKLDPFLIP IKDLNVRPKTIKSLI EKNK WGKDSLFN LAICRKLKLDPFLIP IKDLNVRPKTIKSLI 17982 48350 A 18089 659 826 17983 48351 A 18090 3531 3878 17984 48352 C 18091 1 2793 17985 48353 A 18092 1607 1774 17986 48354 B 18093 1 1666 17987 48355 A 18094 1 1068 17988 48356 B 18095 1 1549 17989 48357 A 18096 557 623 KKWINYWTHTPSQ P 17990 48358 B 18097 1 1231 1231 17991 48359 A 18098 263 1461 GGSSCYPSWEATIR K\RTLKVLTFLVLFT SHTMVATTWPKC LKRLIKHRPCTCTH AWFDERFNQTMQP LEDDTYRWLSLE GFCLQRPFQDKQGT TGRDLTQQQRLQRI DTIKELFRVVFONV VGCRRCAVVGNSG PEIDSHDFVLRMNK DVGTKTTHHLVYP VSMILVPFKTIDLENT TSHTMVATTYPPKCIDLENT TSHTMVATTYPPKCIDLENT TSHTMVATTYPPKCIDLENT TSHTMVATTYPPKCIDLENT TSHTMVATTYPPKCIDLENT TSHTMVATTYPPKCIDLENT TSHTMVATTYPPKCIDLENT TSHTMVATTYPPKCIDLENT TSHTMVATTYPPKCIDLENT TSHTMVATTYPPKCIDLENT TSHTMVATTYPPKCIDLENT TSHTMVATTYPPKCIDLENT TSHTMVATTYPPKCIDLENT TSHTMVATTYPPKCIDLENT TSHTMTYIPVPAKIRV PAFIKYVFDNWLQG GILSVIFSMHVCDENT GILSVIFSMHVCDENT TSHTMTYIPVPAKIRV PAFIKYVFDNWLQG GILSVIFSMHVCDENT TSHTMTYIPVPAKIRV PAFIKYVFDNWLQG GILSVIFSMHVCDENT TSHTMTYIPVPAKIRV PAFIKYVFDNWLQG GILSVIFSMHVCDENT TSHTMTYIPVPAKIRV PAFIKYVFDNWLQG GILSVIFSMHVCDENT TSHTMTYIPVPAKIRV PAFIKYVFDNWLQG GILSVIFSMHVCDENT TSHTMTYIPVPAKIRV PAFIKYVFDNWLQG GILSVIFSMHVCDENT TSHTMTYIPVPAKIRV PAFIKYVFDNWLQG GILSVIFSMHVCDENT TSHTMTYIPVPAKIRV PAFIKYVFDNWLQG GILSVIFSMHVCDENT TSHTMTYIPVPAKIRV PAFIKYVFDNWLQG GILSVIFSMHVCDENT TSHTMTYIPVPAKIRV TSHTMTYIPVPAKIRV TSHTMTYIPVPAKIRV TSHTMTYIPVPAKIRV TSHTMTYIPVPAKIRV TSHTMTYIPVPAKIRV TSHTMTYIPVPAKIRV TSHTMTYIPVPAKIRV TSHTMTYIPVPAKIRV TSHTMTYIPVPAKIRV TSHTMTYIPVPAKIRV TSHTMTYIPVPAKIRV TSHTMTYIPVPAKIRV TSHTMTYIPVPAKIRV TSHTMTYIPVPAKIRV TSHTMTYIPVPAKIRV TSHTMTYIPVPAKIR							
ARIAKTILSKKNIAC LYYKATVTKTAWY QWNRTEASEITPHI							
LYYKATVTKTAWY QWNRTEASEITPHI							`
QWNRTEASEITPHI EKNKKWGKDSLFN LAICRKLKLDFFLIP IKDLNVRPKTIKSLE 17981 48349 B 18088 1 2016 17982 48350 A 18089 659 826 17983 48351 A 18090 3531 3878 17984 48352 C 18091 1 2793 17985 48353 A 18092 1607 1774 17986 48354 B 18093 1 1666 17987 48355 A 18094 1 1068 17988 48356 B 18095 1 1549 17989 48357 A 18096 557 623 KKWINYWTHTPSQ P 17990 48358 B 18097 1 1231 1231 17991 48359 A 18098 263 1461 GGSSCYPSWEATIR KURTLKVLTFLVLFI SHTMIVATTWFPKC LKRLIKHRPCTCTH AWFDERFNQTMQP LEDDTYRWLSLE GFCLQRPFGDKQGT TGRDLTQQQRLQR DTIKELFRVVPGNV VGCRRCAVVGNSG PEIDSHDFVLRMNK DVGTKTTHLVYPI VSMILVPFKTIDLEV TISHTYIPVPAKIRV PAFIKYVFDNWLQC GILSVIFSMHVCDEV GILSVIFSMHVCDEV TISHTYIPVPAKIRV PAFIKYVFDNWLQC GILSVIFSMHVCDEV TISHTYIPVPAKIRV TISHTYIPVPAKIRV PAFIKYVFDNWLQC GILSVIFSMHVCDEV TISHTYIPVPAKIRV							ARIAKTILSKKNIAGGIMLPDFK
EKNKKWGKDSLFN LAICRKLKLDPFLIP IKDLNVRPKTIKSLI			i	İ			LYYKATVTKTAWYWFQNRDID
LAICRKLKLDPFLIP IKDLNVRPKTIKSLE 17981 48349 B 18088 1				į			QWNRTEASEITPHIYNHLIFDKP
IKDLNVRPKTIKSLI 17981 48349 B 18088 1 2016							EKNKKWGKDSLFNKWCWENW
17981			į l	ĺ			LAICRKLKLDPFLIPYTKINSRW
17982			į				IKDLNVRPKTIKSLEENLGIPFR
17983	7981 48	8349	В	18088	1	2016	
17984	7982 4	8350	Α	18089	659	826	
17985	7983 4	8351	Α	18090	3531	3878	
17985	7984 4	8352	C	18091	1	2793	
17986			Α		1607		
17987			$\overline{}$		1	<u></u>	
17988							
17989 48357 A 18096 557 623 KKWINYWTHTPSQ P 17990 48358 B 18097 1 1231 17991 48359 A 18098 263 1461 GGSSCYPSWEATIR K\RTLKVLTFLVLFI SHTM\VATTWFPKQ LKRLIKHRPCTCTH AWFDERFNQTMQP LEDDTYRWWLSLE GFCLQRPFGDKQGT TGRDLTQQRLQRI DTIKELFRVVPGNV VGCRRCAVVGNSG PEIDSHDFVLRMNK DVGTKTTHHLVYPI VSMILVPFKTIDLEV TISHTYIPVPAKIRV PAFIKYVFDNWLQQ GILSVIFSMHVCDEV							
17990 48358 B 18097 1 1231 17991 48359 A 18098 263 1461 GGSSCYPSWEATIR K\RTLKVLTFLVLFI SHTM\VATTWFPKQ LKRLIKHRPCTCTH AWFDERFNQTMQP LEDDTYRWWLSLE GFCLQRPFGDKQGT TGRDLTQQQRLQRI DTIKELFRVVPGNV VGCRRCAVVGNSG PEIDSHDFVLRMNK DVGTKTTHHLVYPI VSMILVPFKTIDLEV TISHTYIPVPAKIRV PAFIKYVFDNWLQQ GILSVIFSMHVCDEV							KKWINYWTHTPSQD*TRKKLN
17990 48358 B 18097 1 1231 17991 48359 A 18098 263 1461 GGSSCYPSWEATIR K\RTLKVLTFLVLFI SHTM\VATTWFPKO LKRLIKHRPCTCTH AWFDERFNQTMQP LEDDTYRWWLSLE GFCLQRPFGDKQGT TGRDLTQQQRLQRI DTIKELFRVVPGNV VGCRRCAVVGNSG PEIDSHDFVLRMNK DVGTKTTHHLVYPI VSMILVPFKTIDLEV TISHTYIPVPAKIRV PAFIKYVFDNWLQO GILSVIFSMHVCDEV	7,707	0337	n l	10070	337	023	,
17991 48359 A 18098 263 1461 GGSSCYPSWEATIR K\RTLKVLTFLVLFI SHTM\VATTWFPKQ LKRLIKHRPCTCTH AWFDERFNQTMQP LEDDTYRWWLSLE GFCLQRPFGDKQGT TGRDLTQQQRLQRI DTIKELFRVVPGNV VGCRRCAVVGNSG PEIDSHDFVLRMNK DVGTKTTHHLVYPI VSMILVPFKTIDLEV TISHTYIPVPAKIRV PAFIKYVFDNWLQQ GILSVIFSMHVCDEV	7000 4	9259	D	18007	1	1221	
K\RTLKVLTFLVLFI SHTM\VATTWFPKQ LKRLIKHRPCTCTH AWFDERFNQTMQP LEDDTYRWWLSLE GFCLQRPFGDKQGT TGRDLTQQQRLQRI DTIKELFRVVPGNV VGCRRCAVVGNSG PEIDSHDFVLRMNK DVGTKTTHHLVYPI VSMILVPFKTIDLEV TISHTYIPVPAKIRV PAFIKYVFDNWLQQ							CCCCVDCWEATIDTAVAVTID
SHTM\VATTWFPKQ LKRLIKHRPCTCTH AWFDERFNQTMQP LEDDTYRWWLSLE GFCLQRPFGDKQGT TGRDLTQQQRLQRI DTIKELFRVVPGNV VGCRRCAVVGNSG PEIDSHDFVLRMNK DVGTKTTHHLVYPI VSMILVPFKTIDLEV TISHTYIPVPAKIRV PAFIKYVFDNWLQQ	7991 40	.8339	A	18098	203	1401]
LKRLIKHRPCTCTH AWFDERFNQTMQP LEDDTYRWWLSLE GFCLQRPFGDKQGT TGRDLTQQQRLQRI DTIKELFRVVPGNV VGCRRCAVVGNSG PEIDSHDFVLRMNK DVGTKTTHHLVYPI VSMILVPFKTIDLEV TISHTYIPVPAKIRV PAFIKYVFDNWLQG				i			
AWFDERFNQTMQP LEDDTYRWWLSLE GFCLQRPFGDKQGT TGRDLTQQQRLQRI DTIKELFRVVPGNV VGCRRCAVVGNSG PEIDSHDFVLRMNK DVGTKTTHHLVYPI VSMILVPFKTIDLEV TISHTYIPVPAKIRV PAFIKYVFDNWLQG		Į		i			SHTM\VATTWFPKQMVLELSEN
LEDDTYRWWLSLE GFCLQRPFGDKQGT TGRDLTQQQRLQRI DTIKELFRVVPGNV VGCRRCAVVGNSG PEIDSHDFVLRMNK DVGTKTTHHLVYPI VSMILVPFKTIDLEV TISHTYIPVPAKIRV PAFIKYVFDNWLQG		[LKRLIKHRPCTCTHCIGQRKLS
GFCLQRPFGDKQGT TGRDLTQQQRLQRI DTIKELFRVVPGNV VGCRRCAVVGNSG PEIDSHDFVLRMNK DVGTKTTHHLVYPI VSMILVPFKTIDLEV TISHTYIPVPAKIRV PAFIKYVFDNWLQG							AWFDERFNQTMQPLLTAQNAL
TGRDLTQQRLQRI DTIKELFRVVPGNV VGCRRCAVVGNSG PEIDSHDFVLRMNK DVGTKTTHHLVYPI VSMILVPFKTIDLEV TISHTYIPVPAKIRV PAFIKYVFDNWLQG		1					LEDDTYRWWLSLESVMFSVTM
DTIKELFRVVPGNV VGCRRCAVVGNSG PEIDSHDFVLRMNK DVGTKTTHHLVYPI VSMILVPFKTIDLEV TISHTYIPVPAKIRV PAFIKYVFDNWLQG				į			GFCLQRPFGDKQGTTQPVYINQ
VGCRRCAVVGNSG PEIDSHDFVLRMNK DVGTKTTHHLVYPI VSMILVPFKTIDLEV TISHTYIPVPAKIRV PAFIKYVFDNWLQC				į			TGRDLTQQQRLQREKKPNNLN
PEIDSHDFVLRMNK DVGTKTTHHLVYPI VSMILVPFKTIDLEV TISHTYIPVPAKIRV PAFIKYVFDNWLQC GILSVIFSMHVCDEV				1			DTIKELFRVVPGNVDPMLEKRS
PEIDSHDFVLRMNK DVGTKTTHHLVYPI VSMILVPFKTIDLEV TISHTYIPVPAKIRV PAFIKYVFDNWLQC GILSVIFSMHVCDEV				1			VGCRRCAVVGNSGNLRESFYG
DVGTKTTHHLVYPI VSMILVPFKTIDLEV TISHTYIPVPAKIRV PAFIKYVFDNWLQO GILSVIFSMHVCDEV							PEIDSHDFVLRMNKAPTAGFEA
VSMILVPFKTIDLEV TISHTYIPVPAKIRV PAFIKYVFDNWLQO GILSVIFSMHVCDEV			, 1				DVGTKTTHHLVYPESFRELGDN
TISHTYIPVPAKIRV PAFIKYVFDNWLQO GILSVIFSMHVCDEV							VSMILVPFKTIDLEWVVSAITTG
PAFIKYVFDNWLQC GILSVIFSMHVCDEV		l					
GILSVIFSMHVCDE		İ			: :		TISHTYIPVPAKIRVKQDKILIYH
1			į				PAFIKYVFDNWLQGHGRYPST
		ļ					GILSVIFSMHVCDEVDLYGFGA
							DSKGNWHHYWENNPSAGAFR
KTGVHDADFESNV		ļ					KTGVHDADFESNVTATLASINK
IRIFKGR							IRIFKGR
17992 48360 B 18099 1 4855	7992 4	8360	В	18099	1	4855	

SEQ ID	SEQ ID NO:		`	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
17993	48361	A	18100	571	4189	FSLKRSSTSLVSWIPRYFILFEAI VNGSSLMIWLSV*AKILVQRA* VLAFHPS*LRFVKDQVVVDMW HYF
17994	48362	A	18101	7449	7626	SFRSRLFSFHVVELF*VSFLILSS SLIALWSERQSVIISVLLYFLRS AFLPTMWSILE
17995	48363	A	18102	63	329	SGQRRKEVPSPTLPAPPFILKKK EWPDPPRSFFSGGHWAKSSCPS DCSSQLMAC*/CKNAKHQKDN SCLI/STHMKGNSERK*NSGYQ GLG
17996	48364	Α	18103	2	618	GAFLLEVATAAGLSAGLTRLGS QVGGAMRRSKADVERYIASVQ GSTPSTLQKSMKGFYFAKLYYE AKEYDLAKKYICTYINVQERDP KAHRFLGLLYELEENTEKAVEC YRRSVELNPTQKDLVLKIAKLL CKNDVTDGRAKYWVERAAKL FPGSPAIYKLKN/DGWVRGQM VPKPDDEGVQVRGRSGRPDAG WRLNRRSSVGGSAW
17997	48365	Α	18104	1	995	
17998	48366	Α	18105	1	513	
17999	48367	Α	18106	62	521	
18000	48368	Α	18107	1	375	
18001	48369	A	18108	76	480	LVKVCHCTYHHFLTGVVSLPR DGGKGAPCWGRARSPKSALLP YLAPGRG*VGTYEVQN/PNGTS PVQSPATDVGRSPALGNQEP/G T*SPGPQIGRGVTGTARPHTGV SRDSWRRPPSFILNAVANIPKGN APIPKE
18002	48370	Α	18109	445	1128	
18003	48371	Α	18110	784	1189	IPLGKLLLKPVGAPQIERFPFQIP VPLCSLSFPSHSPSSSSARS*VRQ ES*LGTPGSAPTGSP/TSHHPSAS SPPHPSAKAAFLSFVCFLCS*PPP SSCHPCGLGPGPDPSQKMMICT VAHLNQSLIFTVVKVHLL
18004	48372	Α	18111	1	2088	
18005	48373	С	18112	169	213	
18006	48374	Α	18113	1145	1229	
18007	48375	Α	18114	1	975	
18008	48376	A_	18115	1	750	
18009	48377	A	18116	288	368	
18010	48378	A	18117	34	168	
18011 18012	48379 48380	A A	18118 18119	2279	1579 2398	SDCSCPGL*GPPGRWHKRRKSG SPHLSTPEGTPALTSQA
18013	48381	Ā	18120	21	190	OLTIDOTT DOTT ALTOYA

SEQ ID	_	1	SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
18014	48382	Α	18121	li	1773 .	
18015	48383	A	18122	18	290	VAGTWPNFLKKVGIVGKYGTR YGASLRKMVKKIEISQHAKYTC SFCGITKMKRRAVGIWHCGSC MKTVAGGAWTYK*V*FLVVFG SVWTT
18016	48384	С	18123	325	486	
18017	48385	Α	18124	1	498	
18018	48386	A	18125	3	306	RQEIDSIHQVGVYCLALVPANT LPKTPLGGIHISQTKQLFLEGSL HPCNILMCPHTCVTNLPKPRQK QP/WCRPCFRDGWESGCWKTY STSCWKGSGTNRRE
18019	48387	Α	18126	1	1197	
18020	48388	A	18127	1	390	NSRVDDFVAPGLSEAGKLLGLE FPERQRLAAAVG/CSPMSGVIS MSAPFFLGKIIDAIYTNPTVDYS DNLTRLCLGLSGVFLCGAAAN AIRVYLMQTSRQRVVKRLRTSL FSSILGQEVAFSDKAGTGELI
18021	48389	Α	18128	1	695	SGFMDHLEEKADLSELVEKEEL GFFQYYRERCHQKVYHPITKPG GSAKDAAPGGGHHQAGPGQG GDEGIRMMFCVSPNRATFVVS VVPLVSIIDVIYGRYLRKLTKVT QDSLAQATQLAEERIGNVK\TV RAFGKEMTEIEKYPSKVDHVM QKARKEAFARAEGVILNEKSFQ GALEFKNVHFAYPARPEVPIFQ DFSLSIPSGSVTALVGPSGSGKS TVLSLLLRLYDPAS
18022	48390	Α	18129	1	2136	
18023	48391	Α	18130	1	822	
18024	48392 48393	A	18131 18132	1		MVNALSLAARIHAEVPADESPE MTEGYEGFYHLASMKGTVERA DMHYIIRDFDRKQFEARKRKM MEIAKKVGKGLHPDCYIELVIE DSYYNMREKVVEHPHILDIAQQ AMRDCDIEPELKPIRGGTDGAQ LSFMGLPCPNLFTGGYNYHA/P SPKFSAVSAMLLASASSAGRCG ESIFSPLAYTSSPIESDSCGPAL* RFLRH*P/EIFLTQPVF/SENEAID IWN\SPTASVSPLMPSRQNAVRE LFLLTVRRPTGTD

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
18026	48394	A	18133	1	1166	LTHWVYHLYGIQRILIHIPSTSQ
						PNRVRGDIPPLQRIVVPVTVIVQ
	1	1		1		PRLLIMLLPWQTVRLVQVMRV
	1	1	ŀ			LLIKLVAPFIILPAPRRIAVFADE
			;			RQRQAPVVAVVKMNFSRRCVF
						RLHLSGQLLHRQPALRKPRPAF
						TAHAVCPDFFLQPVQQPHQFSR
						EHHAVAVLTPFLLERLCQAAPL
						RFFPLAVLCLDPDERREASRLV
						YRLDTCGVSLHRRQPVSVPAIS
						LALPMLAILVVQTIFMALYAIF
						VTWRMMGKNYDAAVLAAGH
						CGFGLGATPTAIANMQAITERF
		l		}		GPSHMAFLVVPMVGAFFIDIVN
			•			ALVIKLYLMLPIFAG*PMKRR*
						K/SPPQQRQMPDTLRLSSVPSPA
						SSVLSGLLPVVGNVREYGVPGA
						ELPGSSPDRAVNPPL
18027	48395	Α	18134	48	314	LVPMTWVIEASYSIMRKKVGE
						HPHFLDIAHQAMRDCDIEPELN
						PIRGGTDGAQLSFMGLPCPNLF
		1				TGGYN\IMVSMSL*LWKVWKK
		<u> </u>				RCR
18028	48396	В	18135	83	3044	
18029	48397	A	18136	1	1080	DAWADAWOWA DWWWDIND
18030	48398	Α	18137	3	979	DAWADAWSKKADVKVENP\D
						TKEKKPEAKK\VDAGGKVKKG
						NLKA\KKPKKGKP\HCS\RNPCP
						CKKEIGRNSRSGMYSRK\AMYK
1						RK\YSAAKSK\VEKKK\KEKVLA
						T\VTKPVG\GDKNRR*PGVVKLP
						QKCPRYYP/SLEDVPRKACLSH
						GKKTPLVQHVEKNCEASIYPRG PF*FILTG\RHRGK\RVGFPESSL
						AKLAYLLC*LGPLVLKSEVPL\R
						RTHQKFCHLPLSTKI\DISN\VKI
						PKHLTGCLLSRRKKLRK\PR\HQ
			1			EG\EIFDTEKEKYEI\TEQRKIDQ KAVDSQILPKIKAIPQLQGYLRS
1			1			VFALTNGIYPHKLVVLNCLKNP
18031	48399	A	18138	858	1418	VEALINGII FIREV VENCERNP
18031	48400	A	18138	97	516	
18032	48400	A	18140	1	219	
10033	40401	I _V	10140	11	217	<u> </u>

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	location of first codon for peptide		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		
18034	48402	Α	18141	110	768	NLFSGCVESTFRKQPAHQRRSR
			į			ARAETMQRRLVQQWSVAVFLL
						SYAVPSCGRSVEGLSRR/LLKP
						MRPKNVHVPTKVTTYKRAVSE
						HQLLHDKGKSIQDLRRRFFLHH
						LIAEIHTAEIRATSEVSPNSKPSP
						NTKNHPVRFGSDDEGRYLTQE
			•			TNKVETYKEQPLKTPGKKKKG
					ļ	KPGKRKEQEKKKRRTRSAWLD
						SGVTGSGLEGDHLSDTSTTSLE
18035	48403	Α	18142	338	661	
18036	48404	A	18143	443	661	
18037	48405	A	18144	474	531	
18038	48406	Α	18145	208	317	
18039	48407	A	18146	1	807	
18040	48408	Α	18147	1	1122	
18041	48409	Α	18148	283	2204	ASPMAPTSLCVPTLLPSLNQTSS
						SSTAPTGKASWGWPMLRLPGL
						TTPWSLSLTLW*SRPTFPTSSPC
						SFVVLASPSTSLKCWPLSEGA
18042	48410	Α	18149	208	1212	
18043	48411	A	18150	202	1677	
18044	48412	A	18151	1	3285	
18045	48413	A	18152	3	2206	
18046	48414	A	18153	1	1041	
18047	48415	A	18154	2	1460	
18048	48416	A	18155	51	792 362	ALICA A WINGUIN DRESA DI CED
18049	48417	Α	18156	31	302	AHGAAWPSHRKDRESADLGSD GQNGQDLDWGGGCRRCC*RC
						DVLCWGAPHTFRHSYAMHML
						YAGIPLKVLOSLLLGOAPCRCG
						LGQNVGTALPLVSGGPMLFP
18050	48418	Α	18157	609	690	DOQUITO I ALLE DA DOGUITALE I
18051	48419	A	18158	1	2055	
18052	48420	A	18159	3	2236	
18053	48421	В	18160	406	454	
18054	48422	A	18161	1	2151	
18055	48423	В	18162	1	1215	
18056	48424	A	18163	1	594	
18057	48425	В	18164	1	1380	
18058	48426	A	18165	441	692	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		l		sequence		
18059	48427	A	18166	2	1311	QHVRSTVTELWTGNFVNSIQTA
						FAAGIGWLITFCSKGGDCLTSQ
						TRLRLSGMLFLNSLDAKEIYLE
		l				VIHNLPDFELLSANTLEDRLAH
						HRWLLFFHFGKNENSNDPELK
		l				KLKTLLKNDHIQVDRFGCSSAP
		İ		:		DICSNLYVFQPSLAVFKGQGTK
	İ					EYEIHHGKKILYDILAFAKESVN
						SHVTTLGPQNFPANDKEPWLV
	-		<u>.</u>			DFFAPWCPPCRALLPELRRASN
						LLYGQLKFGTLDCTVHEGLCN
						MYNIQAYPTTVVFNQSNIHEYE
	1	l				GHHSAEQILEFIKDLMNPSGGS
	1					LTPTTFIELVTQRKHTEVRI/V*F
						HCPGWPPWPRLWPMAKKPRNL
		İ				LP*SSRPLNT*SSLFCSSNQPDLR
		l				GPLLHRPLSLPRRYRGAQVPNP
						DFGHYSVSPSLIHGLHQCSHLFS
						HSPPPRVLRPPRILTTSTGLPPSP
						RRSATDVHTAYPPTAD
18060	48428	A	18167	1	924	
18061	48429	A	18168	10	1254	
18062	48430	Α	18169	1	618	
18063	48431	Α	18170	1	460	MAAPMTPAARPEDYEAALNAA
					•	LADVPELARLLEIDPYLKPYAV
						DFQRRMFWMFLPFFTLAFSPSV
						LVRFHAADKDIPETGQFTKGRG
	1	1				LLDLQFQVAEDASQSWQKLYL
						LPNCSGQPHQKLVPTAQKFYIT
						VEWGTFFSVPGQVVGSEGSQM
	1	l				SVTVEGSEESEDDKMYMTCSS
	1	l				ALLDSFALC**RHT*DWAIYKR
						KRFIGLTVP/MWLRMPRNHGRS
						CISCLIALANHTKNWFPQRRSFI
						SQWNGVLFSQCLAKLWALKEV
						RCQ
18064	48432	Α	18171	1	582	
18065	48433	Α	18172	1	1578	
18066	48434	Α	18173	1	1293	

E (X=Unknown, sible nucleotide nucleotide insertion) NSDCTFKEDNSL GSNQGPRFAGS ALL WTGGHPSK IDGDFEFETYYS DALNLMSVLNP FQNEITDRNVM EFGQGRMTLTEI LAAEESARKGIR QILDTVDIENYIS GALKVHVDEYD S/AISGFFHTYDS QNCAPSPSGEY ILVWRHCDY/PL RATSNTIATMII
NSDCTFKEDNSL GSNQGPRFAGS ALLWTGGHPSK IDGDFEFETYYS QALNLMSVLNP FQNEITDRNVM EFGQGRMTLTEI RAAEESARKGIR QILDTVDIENYIS GALKVHVDEYD S/AISGFFHTYDS QNCAPSPSGEY ILVWRHCDY/PL
GSNQGPRFAGS ALLWTGGHPSK IDGDFEFETYYS QALNLMSVLNP FQNEITDRNVM EFGQGRMTLTEI LAAEESARKGIR QILDTVDIENYIS GALKVHVDEYD S/AISGFFHTYDS QNCAPSPSGEY ILVWRHCDY/PL
GSNQGPRFAGS ALLWTGGHPSK IDGDFEFETYYS QALNLMSVLNP FQNEITDRNVM EFGQGRMTLTEI LAAEESARKGIR QILDTVDIENYIS GALKVHVDEYD S/AISGFFHTYDS QNCAPSPSGEY ILVWRHCDY/PL
GSNQGPRFAGS ALLWTGGHPSK IDGDFEFETYYS QALNLMSVLNP FQNEITDRNVM EFGQGRMTLTEI LAAEESARKGIR QILDTVDIENYIS GALKVHVDEYD S/AISGFFHTYDS QNCAPSPSGEY ILVWRHCDY/PL
ALLWTGGHPSK IDGDFEFETYYS QALNLMSVLNP FQNEITDRNVM EFGQGRMTLTEI RAAEESARKGIR QILDTVDIENYIS GALKVHVDEYD S/AISGFFHTYDS QNCAPSPSGEY ILVWRHCDY/PL
IDGDFEFETYYS QALNLMSVLNP FQNEITDRNVM EFGQGRMTLTEI RAAEESARKGIR QILDTVDIENYIS GALKVHVDEYD S/AISGFFHTYDS QNCAPSPSGEY ILVWRHCDY/PL
QALNLMSVLNP FQNEITDRNVM EFGQGRMTLTEI RAAEESARKGIR QILDTVDIENYIS GALKVHVDEYD S/AISGFFHTYDS QNCAPSPSGEY ILVWRHCDY/PL
FQNEITDRNVM EFGQGRMTLTEI LAAEESARKGIR QILDTVDIENYIS GALKVHVDEYD S/AISGFFHTYDS QNCAPSPSGEY ILVWRHCDY/PL
EFGQGRMTLTEI RAAEESARKGIR QILDTVDIENYIS GALKVHVDEYD S/AISGFFHTYDS QNCAPSPSGEY ILVWRHCDY/PL
QILDTVDIENYIS GALKVHVDEYD S/AISGFFHTYDS QNCAPSPSGEY ILVWRHCDY/PL
GALKVHVDEYD S/AISGFFHTYDS QNCAPSPSGEY ILVWRHCDY/PL
S/AISGFFHTYDS QNCAPSPSGEY ILVWRHCDY/PL
QNCAPSPSGEY LVWRHCDY/PL
LVWRHCDY/PL
RATSNTIATMII
CLYALNCSGDIL
RLMSWLHCLVP
QWQSAKVV*FA
RAIIYQQSRKQ
RSIITVITQSPTK
GDTDAAFGALI
PLPALKPLCVM
WFSRAG*LGEL
PQQQTVSGNG
HSWHY/YTISFQ
PIF*VAKAGRK
IT\D*PVSLSRSQ
GW/QQLDEEIR `
TRDRRKGVQPP
LERRIARLETG
TRHLISNCQISET
YLLYRAQKER
KVRRLEDDTG
AGIISIT
ETAGGAAAHP
IQTQLPGGRGGI
GGGGDDGRRR
CVQRAGPAMDA
RPKDQEVLHRL
RHAGQPAGPA
HLHAGIQPPE/V
RRCPAFSSGTVK
PPLLRKGAASIW

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence		in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
18077	48445	Α	18184	1	2130	
18078	48446	A	18185	2	433	EMTLDSRVAAAGDLFVAVVGH QADGRRYIPQAIAQGVAAIIAE AKDEATDGEIREMHGVPVIYLS QLNERLSALAGRFYHEPSDNLR LVGVTGTNGKTTTTQLLAQWS QLL/GRNQRGNGHRW*RPAGES DPDRKYNRFGSRCSA
18079	48447	A	18186	114	586	SYITALGQFSRSGIQLLDRLGNF GTGWPNFDRKRFGSTMRTTVN GQTR/GPPQRRPAADNLDF**H QRPAGAANRPSGNGHGLGRRA AVTHPSLNQQLPDKRNRHPKRI CCAGKLSALSGTKRSAMKHSM LERVSTKFAVWKMTTGACWN GISSWIKR
18080	48448	В	18187	1	3577	
18081	48449	A	18188	2	861	YHSINAASACSALHPVAFLTILT CRTCSSSLVATTSTMRLPNTLP DLIISPVESILSTSFCAVPAFIREE PVTTSAPTIGVMANSAVVETAL PGLQEIPTVRQPSWRAYSSPAIT YGVRPLAAMP/SHHILCR*INIF QIINTAFPAIFRAFHRVLNGIFSA SNQTNHQTRFHAVCGRTFGDV GWVAKSYQVAEERGSITEHQV QGHQSDDTEHFHGGEIERQQQ HYHRQRGDPVGEIGEPELSISRD RHHVCHTGYNPAEPVSPAGEIS NGTANGNKLNMSVCEIAP
18082	48450	A	18189	560	2021	
18083	48451	A	18190	525	804	GG*CRNWCDHYANGCRDTSGT LYDKLAELGPQGLSPR*NNWQ TARRNQSSGRNSCHLRREVE*R RLDKALRRIRQFSAPDEGAKCP WPGIIDCSNHFARLRCQYGRFL EVTSAGDEKPYPLYDHHVGLV VEQLRCIITAKVRTRHFYYVLK SIHPAVAGLPALRDCGELFNSV YCRKPATGPELQADVMVVVAY GLILRKQCWRCRVLAVSTFMFT AATLARCCTNPTLTRAGDAET GVTIMQMDVEIPVVRCTTSWQ SLAHKAYHHVETTGRRHGETR VQDETLVTYAEKLSKEERVLTG HFRQHSLNAALRFQSMANELA
18084	48452	A	18191	1	266	SQYPLVAGHEVIGRVVALGSA AQDKGLQVGQRVGIGWTARSC G/QLRRLY*R*SDQLRARCGAD DYESRWLCREVACGLAMGDST ARKY

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
18085	48453	Α	18192	5	385	
18086	48454	A	18193	330	598	VMRFAIVVTGPAYGTQQASSAF QFAQALIADGHELSSVFFYREG VYNANQLTSPASDEFDLVRA/S ATTECATWCGAEYLRSGSITPW RC
18087	48455	A	18194	1	2652	
18088	48456	A	18195	1171	2091	
18089	48457	A	18196	486	837	
18090	48458	A	18197		2943	MAIYKQSQHMTEVVRRCPHHE RCSDSDGLAPPQHLIRVEGNLR VEYLDDRNTFRHSVVVPYEPPE VGSDCTTIHYNYMCNSSCMGG MNRRPILTIITLEDSSGNLLGRN SFEVRVCACPGRDRRTEEENLR KKGEPHHELPPGSTKRALPNNT SSSPQPKKKPLDGEYFTLQIRGR ERFEMFRELNEALELKDAQAG KEPGGSRAHSSHLKSKKGQSRL GTRFLCQPAVTVIPVKIVRVTG DTVIRPHAEGAVRVQLRVGQA VTCRVCGIRDADIQIRCGGVNA GQPAGGAVAVTPGLARAADAD EFAVVVPGQPFHGGQAVRYTA VRQPLRVGRMSGLMLAKTCFR LQVMHRLTLTVFSQKVFTESIE VVEALKVNLMRVFLRRMDLRQ HRKMIMIDNYIAYTGSMNMVD PRYFLRDALPHCPNPLTGPTFA VLEKLGYPHTFVSTDFNTVALD VKTFGIKANDDKAKLKALGEIA HACVDATIIYCKSPTVAGLVAR ELIKLGHGTPTENPHVYWSPLE TVLACILARYRGHCSNTLATNS MLENCAFCSAPPRSLKESTLSL NTVVIYDNRDGTRSIDKFNRFCI HAFPKIIRNAIYEVEQHT*RACR YQ*DVGWFNIYQPGISLAASKH CSTSSR*RWI*LDWI/CASASLL DEATAAAEAMAMAKRVSKLK NANRFFVASDVHPQTLDLMSY
18091	48459	A	18198	643	776	(1222.1101
18091	48460	A	18199	1	1257	
18092	48461	A	18200	3	346	
18094	48462	A	18201	1120	1335	
18095	48463	A	18202	312	1108	
18096	48464	A	18203	1	744	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18097	48465	Α	18204	381	1112	GVEPSPPTNPPPATAIPRATCVA
						AARAASPSPS*SHQPHPPPLAYD
		1				RCST*PTPTTTAGQDHDHGRR
						WRSAPAPRSDHLSDMLQE/RSM
						SLLPLILSSTRSGSNPLRSTFLSR
		l				SRSISKNCSAEV*SNNSAGVGTI
]	STS/RSTYDINRADTQVRRAVN
						NYDIIVMSNSFNGQSEHQVWIG
						GQF/ILHQICPLTQFQYGAEPDR
						KE*CQAPASKWALLHK*HRPFP
						QDRKAGECLLHEYEDLYPSVIH
18098	48466	Α	18205	2	307	HPARHKFVKGSSEHRKSPVKPP
	,	1				PIPSH*PP*PENTNPMRPLAPTDP
						VTNLRSLMTGRQRFTPYNNVS
						RSDPSTTARSSSADRVNASEYP
						TSSEETSGRSSR
18099	48467	В	18206	1	723	
18100	48468	С	18207	127	1410	
18101	48469	Α	18208	1	1773	MGKKQSRKTGNSKNQSASPPP
	ļ					KERSSSPAMEQSWTDNDFDEL
						REEGFRRSNY\SSY*EEIRT*WQ
						TSLKALKRN*MMM/ILLASPSR
						TPISGPNGDPAGPKFVKAAATT
						GKARQRFHPATTTFHGRTPAPP
						LAHPAPTVSTPASTQHPARRHQ
						VARPDRSSTPLPARIALRVFCPT
						PSTAPPEHPRQNPNPAPQTDSA LPGADDVRVSTTDPERRHPHPT
						NTTTPGPGRVLRKELNRPRRPI
						HHRRRRFHVQRAWQLPVPHRH
		;				HHLDHTSHTRRRLRMTDVRLD
						RPQPQRPPRITITTVGGDQRLRL
						DRSYQPESPCDGDLRQLQMAG
						RAPCSPLSMLVLEKCDNLKTCH
						TSHGSVMAETAVINHKKRKNS
						PRIVQSNDLTEAAYSLSRDQKR
						MLYLFVDQIRKSDGTLQEHDGI
						CEIHVAKYAEIFGLTSAEASKDI
						RQALKSFAGKEVVFYRPEEDA
						GDEKQNAKNEHGTTSSLPTQFK
						TDHMESQKDHPHKEETSPKVP
						HLVQESQKDHLHKEATNLKVP
						HPHPGKPQGPPPQEGNKPQRPP
						PPGRPQGPPPPGGNPQQPLPPPA
						GKPQGPPPPPQGGRPHRPPQGQ
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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18102	48470	Α	18209	944	1577	RLCCLEISSTGYPKSSISSSKYPQ
1						IPRAGAQCSQAV*SGSNPLRSTF
			1			LSRSRSISKNCSAEV*SNNSAGV
						GTISTS/GSTYDINRADTQVRRA
						VNNYDIIVMCNSFNGQSEHQV
						WIGGQF/ILHQICPLTQFQYGAE
						PDRKE*CQAPASKWALLHK*H
						RPFPQDRKAGECLLHEYEDLVP
						IRDTLRLFPGGRYLPRAKHVAP
						SEPDPTHHAYD
18103	48471	A	18210	3	634	DRVFDSGSKWRSEQDWAAAK
						AVGQCSENGDWRCCGNGRYP
						ARGPAVPA*PSLCLHHPR*RCT
		ŀ				CCMPVYR
18104	48472	Α	18211	3	255	
18105	48473	Α	18212	101	547	TSVELMDLCPIRLCRTFSGIPAY
						SMCIA*EWRNVCGVTGTENVT
						PSAAAAATASPIQVLTVLSVTS
						QIRAFSVLPVRRLRRSMSLSRIN
						TCRTEYPRCAVVCLLWKDAVS
						CLSTTFCARLCGQNTWLPRPCR
						HVNRAASDASVSLSTVT
18106	48474	В	18213	1	1113	
18107	48475	В	18214	220	1173	
18108	48476	Α	18215	1	509	MPNHKAILIKFKKTEIIPITLLDH
						STIKIEINIKKIAQNHTITWKLNN
		=				LLLNYFWPLTVSLCEMCKDFYS
						LGIEFSRFPLQELQNKDCALEK/
						HSHWAQMWP*YPCPAKHWWS
						CRREHVLVVN/DCSTS*NSYWH
						FSD*LRAMKRRNEQCKR*AIRS
						LSRSLPARQDLKLLS
18109	48477	A	18216	3	281	
18110	48478	Α	18217	878	1135	
18111	48479	С	18218	1	1602	
18112	48480	Α	18219	33	608	VDSLRFQSPSCSVYSTDTR*ISE
		l				T**NKL*PPKPVIACPKMWIQV
						GQKFRQKAAATP*KAR*SPRPIP
						SH*QP*PKIQTRCGPQPPQIRSPT
						QKPHDRQTALSHRTTTFHGPTP
						APPLAHPAPTVSTPASTQHPAR
						RHQVARPDRSSTPLPARIALRVF
						CPTPSTAPPEHPRQNPNPAPQTD
		<u>L</u> .			_	SALPGAVRG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18113	48481	A	18220	3	893	NPRKRTEQGKAQNKVQWWLT
		į				RTPVMVQKRPTIVSLKSSPDSPL
e.						ETYSMFLFLGMWVGVGVEM/C
						SVKLEEQVNTYHYQRILLGISS
						DNAILLEEESEEGRPMHISTIKE
						PSPDSPILYLTLNLKLLTSLILFH
						YGESWNLLRADQRLIFAKSWP
						RASRYQQGHQDLFILRSDLPSQ
			İ			VFIRDKLMERRNRRTGRTEKAR
						IWEVTDRTVRTWIGEAVAAAA
		į				ADGVTFSVPVTPHTFRHSYAM
		į.				HMLYAGIPLKVLQSLMGHKSIS
						STEVYTKVFALDVAARHRVQF
						AMPESDAVAMLKQLS
18114	48482	A	18221	1	327	LISLRQRQQRTPGKPGKGPRPIP
						SPWPP*PENTNPMRPPSPTRSGH
						QPRSLMTGRQRFHTAQQRFTV
						RPQHHRSLIQRRPCQRQRVPNI
			}			QRGDIRSLVQIGHQRRCLRG
18115	48483	В	18222	1	1419	
18116	48484	A	18223	1	2784	
18117	48485	Α	18224	936	1157	RTSTGLSGCIRCG\CASEGGQAV
1						FV/WPRGPPETPCSRCPAAEGSP
						G*SPPREDPSSGSPGKRQTCTQT
						SARSPS
18118	48486	В	18225	67	1314	
18119	48487	Α	18226	302	1423	CRWRFRFIMPFVMASMSAECL
						MNYNSTAMSGKTGLQNDTTLI
						ARPWVKPYLGLEIMRIVGN*PH
		l				LTWCFPCTMYIPLFSQLLCCPVS
		l				FGQCGAFRNPGHPPSPNPEFFLG
						PRRCNPSLGHKALVQEAASDTP
						GKS\PGKSPPGPIPKPIGPP*PGK
ļ						HKPDAAPSPHRSGSPNPEAS*PE
<u> </u>		ŀ		-		DSAFTPHNNVFTVRPQHHRSLI
				Ì	İ	QRRPCQRQRVPNIQRGDIRSLRF
						GCFARHHPRHHRNIHDRTRTRP
				1		RRLTRHSRGLFEDDVRVSTTDP
		1				ERRHPHPTNTTTPGPGRVLRKE
				1		LNRPRRPIHLRRRRFHVQRAWQ
						LPVPHRHHHLDHTSHTRRRLR
	:					MTDVRLDRPQPQRPARITITAV
						GGDQRLRLDRITKPRARPN
18120	48488	A	18227	1731	2392	
18121	48489	В	18228	1	1422	
18122	48490	В	18229	298	1464	
18123	48491	В	18230	1	1800	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18124	48492	A	18231	745	1250	HTQNQRSGRSQNDF*RRRKNIS
						ERSAVYR*RQLYIRRQPYAERP
						SLR*RQRP*IPCIRSQHGNRKRIP
						RRRIFI*QSVLRRRHELLP*RKPE
		1				ASAERHKTRC*VSERRPRYHRV
						KYDYH*KSIYSR*AHGAA*PSH
					ł	RKDRESADLGSDGQNGQDLDC
		1				RGGCQPLLLTV
18125	48493	Α	18232	1	1661	MAETAVINHKKRKNSPRIVQSN
						DLTEAAYSLSRDQKRMLYLFD
						DQIRKSDGTLQEHDGICEIHVA
		1				KYAEIFGLTSAEASKDIRQALKS
						YAAKEVVLYRPEEDAGDEEGY
1						ESFPWFIKRAHSPSRGLYSVHIN
						PYLIPFFIGLQNRFTQFRLSETKE
ŀ			-			ITNPYAMRLYESLCQYRKPDGS
į						GIVSLKIDWIIERYQLPQSYQRM
İ		ĺ				PDFRRRFLQCINAGHEMTKAIAI
						AQFNDDSPEARKITRRWRIGEA
						ADLVGVSSQAIRDAEKAGRLPH
						PDMEIRGRVEQRVGYTIEQINH
						MRDVFGTRLRRAEDVFPPVIGV
						AAHKGGVYKTSVSVHLAQDLA
						LKGLRVLLVEGNDPQGTASMY
			ŀ			HGWVPDLHIHAEDTLLPFYLGE
						KDDVTYAIKPTCWPGLDIIPSCL
						ALHRIETELMGKFDEGKLPTDP
						HLMLRLAIETVAHDYDVIVIDS
						APNLGIGTINVVCAADVLIVPTP
						AELFDYTSALQFFDMLRDLLKN
			ł			VDLKGFEPDVFIRDKLMERRNR
						RTGRTEKARIWEVTDRTVRTWI
						GEAVAAAAADGVTFSVPVTPH
						TFRHSYAMHMLYAGIPLKVLQ
						SLMG*YVHCKALWMDCAHV**
		1				TKEKIHSLLHRRHPLQGDKEPL
					1	PSPRNSSMPAVYPYWLPQRSIRI
		<u> </u>				FQH!
18126	48494	A	18233	162	446	
18127	48495	B	18234	144	2537	
18128	48496	C	18235	1	873	DEECTDEEDWA DWOJOL DAWS
18129	48497	A	18236	700	1198	PFFGTDFFPWARWCIQLRHKFW
						SKGKQRTTKKAPVKPPGPSPAI
						DHP*PGKHKPDAAPSPHRSGHQ
						PKTSMTGRQRFHTAQQPFHGPT
						PAPPLAHPAPTVSTPASTQHPA
						RRHQVARPDRSSTPLPARIALR
		1				VFCPTPSTAPPEHPRQNPNPAPQ
	<u></u>					TDSALPGAVRG

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
18130	48498	Α	18237	363	940	LQTVLQGYSNENSMFVKQKGS
10.50	10120	-	1020			PPGKPGKKKPCFWGAPKNGAD
						IPSQSWRPSGNPGTTPPLPKPPSF
						FPGAPRMGQSQAWGHKGFWF
						KKGKPTEPPGKKARVKKPPGG
		1				QSPKAIGPTRGPGKPPKPDMRA
		ļ				PKPPTDSGHQTQKPS*PADSAFT
						PHINVSRSDPRTTARSSSADRVN
		1				ASEYPTSSEETSGRSSR
18131	48499	Α	18238	861	1517	SLRPHLSGFYQARTL*DQRS*A
		ļ				DHE/DISKNCSAEV*SNNSAGV
		1				GTISTS/RSTYDINRADTQVRRA
		l				VNNYDIIVMSNSFNGQSEHQV
		ļ				WIGGQF/ILHQICPLTQFQYGAE
						PDRKE*CQAPASKWALLHK*H
						RPFPQDRKAGECLLHEYEDLVP
						IRDTLRLFPGGRYLPRAKHVAP
						SEPDPEQDEQKLRFCKRHLYGQ
]					QPRSPVEIRLQHVAIAYQTHHA
18132	48500	В	18239	1014	2337	
18133	48501	Α	18240	1495	1947	IVTLCFPSYCLHCEAGP/LPCGH
						PWLWLC*VLGSPG*TQRTGSW
						AMPGVQASKPACG*KVAPAAR
	<u> </u>					RVPHPLHMLPHSGHSAASGGA
						SCPLCRG*ATGSDQVLKHAFRK
						GRSSNCASNGTPPPNADFLGQD
						VHPARHKLRQKSASEPRKKAR
18134	48502	В	18241	1	2055	
18135	48503	A	18242	1	909	
18136	48504	В	18243	184	395	
18137	48505	Α	18244	447	667	HRPRRRFQIKPQSQVPEHAPLPL
						PTASDPP/AHSPRCPSESRLSPGG
						QRLGFPSSVLNCSKCSWQLAAC
		<u> </u>				MGFSF
18138	48506	A	18245	265	890	
18139	48507	В	18246	1	551	
18140	48508	Α	18247	407	636	RIMVHATGLMKHASSPGCWDL
						NRRTRRCGVRRVTE*QRAS*KR
		ł				*/ERDYISIMPKPDGLTAAKNLA
						EAFEHYNECIRI
18141	48509	A	18248	1	1448	
18142	48510	Α	18249	817	930	
18143	48511	В	18250	303	461	
18144	48512	Α	18251	612	3359	
18145	48513	A	18252	1	462	

NO: of peptid sequence 18146 48514 18147 48515		in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion) RGGIEYLLLVVFILTRTRFIQAV EHRRNGRRVAEKIVPELITHFM AEIRDLLQRTGVAL/W*HQSAA YLPG*QICALA**IHPGVAAGEY
18146 48514	A		sequence		RGGIEYLLLVVFILTRTRFIQAV EHRRNGRRVAEKIVPELITHFM AEIRDLLQRTGVAL/W*HQSAA
		18253	<u> </u>	749	EHRRNGRRVAEKIVPELITHFM AEIRDLLQRTGVAL/W*HQSAA
		18253	268	749	EHRRNGRRVAEKIVPELITHFM AEIRDLLQRTGVAL/W*HQSAA
					EHRRNGRRVAEKIVPELITHFM AEIRDLLQRTGVAL/W*HQSAA
18147 48515	A				AEIRDLLQRTGVAL/W*HQSAA
18147 48515	A				1
18147 48515	A				
18147 48515	A			1	QRRSIQENGGKVQESGQTAS*M
18147 48515	A		l .		RG*MSVESRSGIAARCPESDAQ
18147 48515	A		r		SGVVHRSRWNQSLPARFGAGR
18147 48515	A				DYREAS
		18254	1793	2193	TQRIPAAQKQGSRCIVDSSAEA
					LSAALAIGNIELVKPSRTMS/DK
1					AAQEIVNSGKAKRVVVSLGPQ
				•	GALGVDSENCIQVVPPPVKSQS
					TVGAGDSMVGAMTLKLAENA
					SLEEMVRFGVAAGSAATLNQG
					TRLCSH
18148 48516	В	18255	1	1557	
18149 48517	A	18256	606	1123	QRDNPSGTDGGRNRAFGAGNI
					HTRGAAQAVERLVDSFPAQEK
					DPGESTGRTDSLPSSRPQKVAY
					PVHRNDRR/ERLWFAAPSSPSN
					ATWGNAEWTLNFSSSDSVTGA
					SSER*NLLFRLATVRCTLPSSVS
					ADGAIVAALATHIGEAELTLES
					QCGAWRATFSSTCSSLPLNPN
18150 48518	Α	18257	110	318	TALKLPLPMSTKVTRCLSNLLK
					PKFNLPPLVQPPKKMLPRHLSR
1					*APNTKTSRVLKKN*PIPIPTRR
10151 40510		100.50	2	3062	KT
18151 48519	A	18258 18259	4435	4954	
18152 48520	A		· · · · · · · · · · · · · · · · · · ·	1959	
18153 48521 18154 48522	A A	18260 18261	2	715	
18155 48523	A A	18262	1	756	MIRRVSVASGNEYPMRPASYIR
18155 48525	TA .	10202	'	750	TAPTVYPSSSPAGTWFAGNAPA
					SSCGFRDRYLIVCSHQTPHVPG
					KREEVNYETGIYWLRHYGYTD
	ı				YSRNAKQRLQRNNLAPILFTTT
				}	VPLGGRQYNNSQHLPQRQSLE
					RIRHHTQAFTALIPHGAAGQGC
					GSGDLHYLSLEDVAVSSKATPD
					GSGVTRKASANLRVTIQIGGFQ
					LRTCGFGPSCRGCHPALQISRAL
					AVAAHQQAADNNMEFANYGP
					FELLQ\ARQLIES
18156 48524	A	18263	1	799	
18157 48525	A	18264	1	270	
18158 48526	A	18265	2	188	LGFFFQRPLGFGQELSKL*EQVL
					LAPEEVGHLGEHLLFRHASEWP
					APRVQLLLLVRVQDLQE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:		li de la companya de la companya de la companya de la companya de la companya de la companya de la companya de	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
18159	48527	A	18266	2791	3274	SKIFPRHDDLLLWRSHRNGDDL RTAGMNAFKQRLTLLGFKITIL TANDYLFADGQ/RSVRKKYPQ AQVSRVAGADLWEELMARAG KE/GTPVFLVGGEPEVLAQTEA KLVRSSSATSAIAWPLLIPCAISS WIRSTAFCCCSWVSEGKPADRS PAPSRTWLRC
18160	48528	A	18267	202	627	
18161	48529	A	18268	1	2390	
18162	48530	Α	18269	1081	1911	
18163	48531	Α	18270	86	272	
18164	48532	A	18271	674	2808	FILKLRTRRR*RHRLIRPWELFQ TARNAEVMKA/LVRHLWLEVK GVDLGRFPERRYGSDKPDLRNP MELTDVADLLKSVEFAVFAGP ANDPKGRVAALRVPGGASLTR KQIDEYGNFVKIYGAKGLAYIK VNERAKGLEGINSPVAKFLNAE IIEDILDRTAAQDGDMIFFGADN KKIVADAMGALRLKVGKDLGL TDESKWAPLWVIDFPMFEDDG EGGLTAMHHPFTSPKDMTAAE LKAAPENAVATLGDPLESPDDG LHNRISTHGCFYVLCGISTNIGV ESTARNAWELGFNLVIAEDACS AASAEQHNNSINHIYPRIARRG RDPQRVMIYIGLPQWSHPKWV RLGITSLEEYARHFNCVEGNTT LYALPKPEVVLRWREQTTDDF RFCFKFPATISHQAALRHCDDL VTEFLTRMSPLAPRIGQYWLQL PATFGPRELPALWHFLDSLPGE FNYGVEVRHPQFFAKGEEEQTL NRGLHQRGVNRVILDSHPVHA ARPHRKLGGTAVWEWEPEGA HWVGDEPGWFSSGDLSKAEY WAQKAADSGDADACALLAQIK ITNPVSLDYPQAKVLAEKAAQA GSKEGEVTLAHILVNTQAGKPD YPKAISLLENASEDLENDSAVD AQMLLGLIYANGVGIKADDDK ATWYFKRSSAISRTGYSEYWA GMMFLNGEEGFIEKNKQKALH
18165	48533	A	18272	532	2103	
18166	48534	В	18273	1	4401	
18167	48535	Α	18274	343	765	
18168	48536	Α	18275	447	709	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			ŀ	sequence		
18169	48537	A	18276	11	1571	MIEVKHLKTLQALRNCGSLAA
10102	10557	1.,	10270	1		AAATLHQTQSALSHQFSDLEQR
						LGFRLFVRKSQPLRFTPQGEILL
		ŀ				QLANQVLPQISQALQACNEPQQ
						TRLRIAIECHSCIQWLTPALENF
						HKNWPQYSVIFKSGVTFDPHFF
						FQQGELDPERAFDDFGIKPMAS
						ASIAQVHSARFNSNGHEVVIKV
		ľ				IRAETLPVIKADLKLIYRLARW
				:		DAAFAAGWSPPAANLSCAGST
			:			KIHRLMNRTCCRNLPTPFSFGA
						KPGLYLVEAMVGGYRATTVVF
						VSDTVALSKVSGIELLVWGAG
						KKQGEANAGSEILWTDGLGVM
						TRGVTDDSGTLQLQHISPERSYI
						LGTLAMVQLREDMAKRQAAGI
						IAPRSGRRMHRTQHKPPMTTE
ł						MGPRAVLREDTIPFLEALKASG
	:					KQRILLTNAHPHNLAVKLEHTG
,		1		1		LDAHLDLLLSTHTFGYPKEDQR
				1		LWHAVAEATGLKAERTLFIDDS
						EGDAMKEKPAVEVRLDKWLW
						AAR/SL*NPRAGP*ND*RR*GAL
						QRAAHEAEQNRRAECHAHSAP
18170	48538	A	18277	2080	2274	RFQVVYFDHEKPSLRHE*FALE
18170	46536	^	16277	2000	22/4	EYTGTRN*GCVNLLDR*KPFPF
						KFMERWLAFSPPGLTLESSRG
18171	48539	В	18278	1	3126	RI WERWEATST GETEESSRG
18172	48540	A	18279	1	341	MLDRLESEILADRVSEESRRWL
	1.02.10					ASCGLTVEQMQNQMDPVYTPA
						RKIHLYHCDHRGLPLALISKEG
						TTEWCAEYD/GMGQPAE*REPA
						SAAAAYPPAGAAV**GVRPVL
						OPPPLL
18173	48541	A	18280	945	1484	
18174	48542	Α	18281	965	1403	
18175	48543	A	18282	703	1149	
18176	48544	Α	18283	161	476	LLPWRSCILRSDPLAAHPVI/LV
						P*ACIFIARNCYPRPSCDSACKL
						RRESCG*FG*TRMVSAQATFSIQ
						ITSA/LPFCEHLPGKRSGDFDTG
				_		EDNALSGQRPEFRHG
18177	48545	A	18284	1	443	MLSEPALRELATAIAGVRKGEN
						DIAVENIIGANIFNIVIVLGLPALI
						TPGEIDPLAYSRDYSVMLLQAG
						KEVLAIERECLAELDQYINQNF
						TLACEKMFWCKGKVVVMGMG
						KS/RAILGEKWPRNVCQHRIAN
					}	HENRDSIRKICCYHR
18178	48546	A	18285	3	550	
					1	· · · · · · · · · · · · · · · · · · ·

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
18179	48547	A	18286	2157	2504	LVSLAGAAYRSPQDPVPYMPIS PGLDWVRQIVCCVMASDAD** RVPRQNALLLPVCTSGPYHALA TGCGFRRGQMLKTTVLPRHYK LYLGASSGMRYSTPATALMPLP LPHLAER
18180	48548	Α	18287	1	621	
18181	48549	Α	18288	1	423	
18182	48550	В	18289	1	1551	
18183	48551	Α	18290	1171	2757	
18184	48552	Α	18291	347	690	
18185	48553	В	18292	1	528	
18186	48554	Α	18293	1	1417	
18187	48555	A	18294	292	699	
18188	48556	Α	18295	859	1377	
18189	48557	Α	18296	1	3036	
18190	48558	Α	18297	1	942	
18191	48559	Α	18298 18299	1955	2124	GGISIDRSGDH*GRGAVYHVAG
			10200			R*NAGRTAELYFASRPTCG/RW LVAAVQGAATARISTIEMSVW AQL*QMCRGKIRAT/ISLPEATA QVISSGSGATKVEDGLGDLNKP VSNQNLVTGIDTPVYNAPSAGS APFGVLADNMRYPILHKLKDR LNQTWYQIRIGDRLAYISALDA QPDNGLSVLTYHHILRDEENTR FRHTSTTTSVRAFNNQMAWLR DRGYATLSMVQLEGYVKNKIN LPARAVVITFDDGLKSVSRYAY PVLKQYGMKATAFIVTTRIKRV
18193	48561	A	18300	1	1191	ADVIVETURE WEREOUTH VIII
18194	48562	A	18301	1	298	MPVVFTVFFLWFPSGLVLYYIV SNLPRYAEYLTVKDADGSVLD QGIALWFP\GPNSFTGEDVLELQ GHGGPVILDLLLIRILTVCGLRI ARPACSPRSL
18195	48563	A	18302	1	1011	4 - 1
18196	48564	A	18303	1	1845	MEDVIDIVACADA VODESTERIO
18197	48565	A	18304		755	MSPVIDKMYVDAYQPFSEEKIS MQEALEKGAQPLREFMLRQTR EADLGLFARLANTGPLQGPEAV PMRILLPAYVTSELKTAFQIGFT IFIPFLIIDLVIASVLMALGMMM VPPATIALPFKLMLFVLVDGWQ LLVALVTGLIISILQAATQINEM TLSFIPKIIAVFIAIIIAGPWMLNL LLDYVRTLFTNLPYIIGRTMLQ VTSEQ/CAILVKPVLLAVTARA GADLHRADSERTQRTKTGGGF CRYWAG

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
18198	48566	Α	18305	1	1977	
18199	48567	Α	18306	1	630	
18200	48568	A	18307	1715	1996	LFRLLKTLTGRRKRSNKSWITT CRGFWGTWCAGWSRP*VVQK CRIYTMWR*WQHAQR*VSPAT LNGLMLITISRKNPHTRKGAGK HCPFSGPS
18201	48569	A	18308	517	675	
18202	48570	Α	18309	480	626	
18203	48571	Ā	18310	757	1146	
18204	48572	A	18311	318	724	
18205	48573	Α	18312	1	1743	
18206	48574	Α	18313	1	1197	
18207	48575		18314	161	613	LLRQGFIVISAGGKRVEGQIKLV FPAKFEAGFRHGVIANLRLYPF PEEKIGEMATAPLSTSGDCRT* YALIGIQ/PTDGDARGSPRYKPA LALLPACVGSAKSTPHGRTPIPA IYVPSSPT*FRALASDSVAQT*Y CRPASQTRRASPARSL
18208	48576	A	18315	177	425	SRYFSATSTWPLVMSIPVTCPFS PASWLST*QSLPLPLPRSKIQQF CRCSGITRPQP*YPDPKSSSFVD AQGSPDHSHNILGLLLRVHFAR LPECAWGPAWLW
18209	48577	A	18316		576	ADSSKKIADIISVIDGIAFQTNIL ALNAAVEAARAGEQGRGFAVV AGEVRNLASRSAQAAKEIKALI EDSVSRVDTGSVLVESAGETM NNIVNAVTRVTDIMGEIASASD EQSRGIDQVALAVSEMDRVTQ QNASLVQESAAAAAALEEQAS RLTQAV\PRSVWQPAHSPINRK HHPVLPIATGALMPGCGS
18210	48578	A	18317	121	1819	
18211	48579	A	18318	1	2151	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
18212	48580	A	18319	3	1127	 GTPYASQPVIRPVEGSKPLIKQV
						LDIGAHGTDKTIAQKLARVAR
						MHFARQRLAAVATFACSAQNA
		İ				IALMEEIAANFSYEMIRLTECIL
		İ				GFTWNRLYQGINVHNAERVRQ
						LAHDGHELVYVPCHRSHMDYL
						LLSYVLYHQGLVPPHIAAGINL
		ĺ				NFWPAGPIFRRLGAFFIRRTFKG
						NKLYSTVFREYLGELFSRGYSV
						EYFVEGGRSR/TGRLLDPKTG\M
						LRGGTRPITLIPIYIGYEHVMEV
		1				GTYAKELRGATKEKESLPQML
		1				RGLSKLRNLGQGYVNFGEPMP
		1				LMTYLNQHVPDWRESIDPIEAV
						RPAWLTPTVNNIAADLMVRIN
						NAGAANAMNLCCTALLASRQR
	<u> </u>	 				SLTREQLTEQLNCYLDLMRNVP
18213	48581	A	18320	1	2470	MFQPSDSGKSFIFNMSVGYNLE
						GIKQPPMQQFIDNMMDASDHP
						KFAQYRDTLNKLLQDDAFLAR
						HGLQEKRESLQALPARIPTSMV
		1				HGVTLSTMHGCPPHEIEAICRY
						MLEEKGLNTFVKLNPTLLGYA RVREILDVCGFGYIGLKEESFD
		1				HDLKLTQALEMLERLMALAKE
		1				KSLGFGVKLTNTLGTINNKGAL
						PGDSTSGVLRTYERLLTAWML
						TCPCADEHQRGESVKLATALT
		1				WWLRGSQNRDEITKDGFQSNH
						AGGILGGISSGQQIIAHMALKPT
		1				SSIDVPGRTINRFGEEVEMITKG
						RHDPCVGIRAVPIAEAMLAIVL
						MDHLLRQRAQNADVKTDIPRC
						AQSIGSFSNGCIVGADTLPIQSE
						HYQVMRTDQRRYFGHPDLVM
		-			·	FIQRLSSQVSNLGMGTVLIGDM
		1				GMPAGGRFNGGHASHQTGLDV
		1				DIFLQLPKTRWTSAQLLRPQAL
		1				DLVSRDGKHVVSTLWKPEIFSL
		1				IKLAAQDKDVTRIFVNPAIKQQ
						LCLDAGTIATGCAKCDPGVSIG
		1				AYAYLLTLRAQCLAHDLPDPLE
						PLEIDGTLLPRYVFIHGGPRVFT
		1	}		1	YYTPKEESIKLFHDYLDLHRSN
		1				PNLDVQMVPVSVMFGRAPGRE
						KGEVNPPLRMLNGVQKFFAVL
						WLGRDSFVRFSPSVSLRRMADE
						HGTDKTIAQKL\ARVARM\HFA
						RQRLAAVGPRLPARQDLFNKL
18214	48582	В	18321	1	1460	
18215	48583	Α	18322	2	234	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
	of peptide		in USSN	location of first	.	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18216	48584	A	18323	282	422	
	48585	Α	18324	11	1974	
	48586	Α	18325	154	1944	
	48587	A	18326	18	386	KACEAVF*DQGLPLGPKVQGQ
						WDWFPGPGALRHQQSVAHQA
						GGFRSLWGYSGVSGQCRRHSD
						*SRPGREARRRRSVAG**SHAL
			<u>.</u>	:		HRQTALFGARSDADLPAAAPR
						YLLYRGLSAAHFRPQAG
18220	48588	Α	18327	181	655	KARSPEDQGFRPRVHIDGGVW
						HLDVGSSHPGAEVGPKGMAVR
						HLKWYASWV*NVVRQFGPYLP
]			WALEN/WRGAAPSTRGHGVC/G
						ITVFGKSDGTT/SHDQY*TAPYG
					:	SPHPPGRWSLRHRYPAHLTSSA
						LAFSDARRNAARQPRRWNPLS
						GYSQVICLPAW
18221	48589	A	18328	1	658	MDDGTSIAPDGDLWAFIGDELK
						MGIPENDRIREQKQKYLRNKSY
į						LHDVTLRAEPYMYWIAGQVKK
						RNMPMELVLLPIVESAFDPHAT
						SGANAAGIWQIIPSTGRNYGLK
						QTRNYDARRDVVASTTAALNM
						MQRLNKMFDGDWLLTVAAYN
						SGEGRVMKAIKTNKARGKSTD
						FWSLPLPQETKQYVPKMLALS
		ŀ				DILKNSKRYGVRLPTTDESRAL
						ARVHLISPVKCARFSRFVFCPA
						VCRDSGGADERRTDSDGCQYA
		l				GGVKPESVMGAEIVEGFSHISV
						TSAWSRGDNSPAPMAFSRLWL
						TVFSMVFKRRETVRSLSAVGA
			ļ			GELRGAAPSTRGPEWTHHWCS
						GCHANGTARSAVVPPDPMPNS
						EVKRRSADGSVGSP*RLITAAK
			 			VGS*RQLKRTKRVGNPRTSGRY
		İ				RCRRKRSSTCLKCWH*VIFSKT
						ASVMAYVCQRPMKAVLWRVC
18222	48590	Α	18329	642	1343	LCLHLPLPTRFPRLQL*EEGGPV
						QPAAMPQWRTLPGPGPRPALPL
						PRRLRGSSLRARPGRLRGPRLR*
						RRHVCGGRRRAPLLLRAGLRRP
		1				RLPRARGPVRRAPLCSRRPLLR
		1		-		PLLRPRLRLRSRLHGSAV*VPSA
		1				PRRKKLARGPAGPQARGPSAL
						PFASGSGTARGRGRGRRCALA
						GPRAPPWPLPGCWVSLAGWDP
	•		l .	ı	I	
			·			GAVSPRTPGCTOOPKDAGGFR
						GAVSPRTPGCTQQPKDAGGFR GWSEAGVHDAQTV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence		in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	lugar-	<u> </u>	1.000	1006	lara	LA DIVOLA PROGRAMO
18224	48592	Α	18331	236	717	ALPVGAAFSSTTGFGSGFGAGG
						ASTFGLGGSTGFGGGATGAGS
						GSNSVLLAAIVAGSSDGAAASP
						ARTSSAAPSGGCD/QERRIVDFK
						RNKDGIPAVVERLEYDPNRSAN IALVLYKDGERRYILAPKGLKA
						GDQIQSGVDAAIKPGNTRLSYG
			ļ			QHREAWR
18225	48593	A	18332	49	291	
18226	48594	Α	18333	75	839	
18227	48595	A	18334	1	579	
18228	48596	Α	18335	223	563	
18229	48597	A	18336	1	1219	
18230	48598	Α	18337	157	864	RGNRQLQQFPRHWRSNRLDGL
						LLQIQLQILGGHTVERHLVVQI
					}	ALVVNGHPGHADGMWILCIRV
			,			LLSLQQNGFRRVPSIGHWTGH*
						AATDQARLMLGGHINAQFAGI QRLEDLGTQLLHTTNGLVEVE
						ARRQILQLGLPLFLLLNQTDVW
						AIPGVHVLRFDVVGRDPS/EVLI
						DPRQNAGLLWKVFATFRPEPTR
						PAVCCPRGGQSGDDRRAGCAC
						NHVELVARHRRSCPWHSDRH
18231	48599	A	18338	1	219	
18232	48600	Α	18339	1	2226	
18233	48601	Α	18340	46	228	
18234	48602	Α	18341	382	1431	
18235	48603	В	18342	1	2400	
18236	48604	В	18343	1	912	
18237	48605	A	18344	2267	2539	
18238 18239	48606 48607	B A	18345 18346	1797 1842	3411 2560	
18240	48608	A	18347	1	1284	
18241	48609	A	18348	1	792	
18242	48610	A	18349	1	978	
18243	48611	A	18350	757	1011	LFGRFSNGFLIVVCQRLRQAVV
		ı				NEQFLPETATLTHRGELLFHSP
		ļ				A*LPGVFRYLAQM*RHAVIQVF
		1				RATGVPEPERCRQRRPHER
18244	48612	Α	18351	148	679	RMSDDKFDAIVVGAGVAGSVA
						ALVMARAGLDVLVIERGDSAG
						CKNMT\GGRLYAHTLEAIIPGFA
						VSAPVERKVTREKISFLTEESAV
						TLDFHREQPDVPQHASYTVLRN
						RLDPWLMEQAEQAGAQFIPGV
						RVDALVREGNKVTGVQAGDDI
1		1				LEANVVILADGVNSMLGRSLG
10045	10013		10252	1170	2241	MVPAS
18245 18246	48613	A	18352 18353	1168	2241 1344	
10240	48614	Α	10333	11	L1344	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
18247	48615	A	18354	440	886	
18248	48616	A	18355	1	2876	MAGVPGFEPGNAGIKNRCLTA
		Ì				WRYPIRTTLSGEWCGRRDLNSQ
		i				EATLQQPLLQAIDLKKHYPVKK
						GMFAPERLVKALDGVSFNLER
l		l				GKTLAVVGESGCGVLACTLFA
						WNRNYFVLLFVGVFLSSFGSTA
						NPQMFALAREHADKTGREAV
						MFSSFLRAQVSLAWVIGPPLAY
						ALAMGFSFTVMYLSAAVAFIV
						CGVMVWLFLPSMRKELPLATG
						TIEAPRRNRRDTLLLFVICTLM
						WGSNSLYIINMPLFIINELH
18249	48617	Α	18356	606	726	
18250	48618	Α	18357	1271	1726	
18251	48619	В	18358	1	1633	
18252	48620	Α	18359	1	407	
18253	48621	Α	18360	427	2307	
18254	48622	Α	18361	120	694	
18255	48623	Α	18362	804	1087	KRQHLCTHSGAIRLRIAGVGSW
						KG/GRPNWRKMAVRWRVPASA
				İ		VVSGWLRRWHCWPVAMPSAE
						SPRIRCGAGCCICRLPLFRCGCC
100.56	10.50.1	ļ.,	100.60	216	100	TPARLTISH
18256	48624	A	18363	216	436	EDDUNUUC CCCTTDI CHOLTD+
18257	48625	A	18364	476	577	FRRWYWGCCSTTRLSHCLTR*
10250	10000	ļ	10265	1150	2744	M*CMRWSLAVCC
18258	48626	A	18365 18366	1159 169	743	
18260	48627 48628	B A	18367	2811	3513	SLPRGWFQPEYGNLEIMEKTYN
18200	40020	A	10307	2011	3313	PQDIEQPLYEHLGFPRVGLRRE
						LKKAQESYWAGNSTREELLAV
					:	GRELRARH\WDQQKQAGIDLLP
						VGDFAWYDHVLTTSLLLGNVP
			-			ARHQNKDGSVDIDTLFRIGRGR
						APTGEPAAAAEMTKWFNTNYH
		İ				YMVPEFVKGQQFKLTWTQLLD
				1		EVDEALALGHKVKPVLAGPGT
		1				WLWLGKVKSEQFYRLSLLNDI
						LAGYQQGLAETGVGPDDIVTA
	1		L	L	l	LAG TQQGLAETGVGPDDIV IA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
18261	48629	A	18368	<u> </u>	2810	 MKQTVYIASPESQQIHVWNLN
10201	140027	``	10500	ī	2010	HEGALTLTQVVDVPGQVQPMV
						VSPDKRYLYVGVRPEFRVLAY
						RIAPDDGALTFAAESALPGSPT
						HISTDHQGQFVFVGSYNAGNVS
						VTRLEDGLPVGVVHVVDGLDR
						LARQLIKNLATLKAGNARVVN
						SNAMSFLAQKGTPHNIVFVDPP
						FRRGLLEETINLLEDNGWLADE
						ALIYVESEVENGLPTVPANWSL
						HREKVAAFAGAEPYHYLNHTL
						GFPRVGLRRELKKAQDSYW
18262	48630	A	18369	2279	2563	MNFGASLNSPTPRSYLSLGGEK
10202	10050	-	10307			KAPVWEHTLGVPSRRPAAGKL
						KKAPERYWGGEHPPVKKWLG
			<u>.</u>			G*ARKCRPRPWETL*KSHLRRR
						PSLGPWAGGL
18263	48631	Α	18370	481	1020	
18264	48632	A	18371	1	998	MNENFGTQICVNNRLSFTDCSG
						YVTRYDHDRFGQVTAVHREEG
						LSQYRAYDSRGQLIAVKDTQG
			i			HETRYEYNAAGDLTTVIAPDGS
						RNGTQYDAWGKAICTTQGGLT
				İ		RSMEYDAAGRVIRLTSENGSHT
						TFRYDVLDRLIQETGFDGRTQR
		ĺ				YHHDLTGKLIRSEDEGLVTHW
						HYDEADRLTHRTVNGETAERW
						QYDERGWLTDISHISERHRVTV
		l				HYGYDSKGRLASEHLTVHHPQ
						TNELLWQHETRLAYKRTGLAN
						RCIPDTCPPVNAYLGTACCQHE
						ICDTPWLSYRDPFTGKSWLSGR
		1	}			YELHKRIYGITAKERKDVDVW
			:			HPDVRFFELYDENNELRGSFYL
						DLYARENKRGGAWMDDCVGQ
						MRKADGSLQKPVAYLTCNFNR
						PVNGKPALFTHDEVITLFHEFG
		1				HGLHHMLTRIETAGVSGISGVP
		1				WDAVELPSQFMENWCWEPEA
		1			•	LAFISGHYETGEPLPKELLDKM
		1				LAAKNYQAALFILRQLEFGLFD
		1				FRLHAEFRPDQGAKILETLAEIK
		1				KLVAVVPSPSWGRFPHAFSHIF
		1		1		AGGYAAGYYSYLWADVLAAD
				1		AFSRFEEEGIFNRETGQSFLDNI
1						LSRGGSEEPMDLFKRFRGREPQ
		1				L*HTVVELPRPFHREILAFRPL*I
						TQAYLRHHR*RA*RC*CLASGC

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence		in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
18265	48633	A	18372	409	533	TGNDAPVPAQSAQYRYLQ*SV RNFCRLSFRLGINLSLPYPHVDD VIPLMAEGKILPYLDIPLQHASP RILKLMKRPG
18266	48634	A	18373	222	659	SAGSVKTQRSGANKQRRTRSRL ASFC/ETLSSVSFPAWRKISFLIP VGLPISNNYVCSLNPVWCAMR LNMPPMSKSICWQKHWKSTVS RWITTRHSFVQTLISTACWRRSP VTQSSWRSTFACSTGLLSHAPP LPIRHCTNITNVWY
18267	48635	В	18374	1	1527	
18268	48636	Α	18375	1	1005	
18269	48637	A	18376	2	314	WVSYCSGSTTL/LLRALQECQR QQGPWASAGDPGEGAGERERL PQALASPPLQVKPARQ/RDPGL RTPDPGDTASPALNGHIFALPA GTAGDCFPRPPRGRKNSGT
18270	48638	Α	18377	3	576	
18271	48639	A	18378	189	563	QGRGSSRPGCGHLDPGGEEWL PQRPGLQHLC\PSAYILIFAGVL VMVTGFLGFGAILWERKGCLS TTTGDRFWKDRQPWTCR/WQT SHRNQSGAGGQWDRFRSHQGH RDHEGRTSNWGSGFPREVA
18272	48640	Α	18379	1	531	
18273	48641	A	18380	1385	1673	AASTDRAAETG**KTR/HRWFL SVKLRTSTPTAVSPKK*KRLAW TRCMTPPSKR*KKRVITPSSSPT SLTSTLPGATVATSPVMPRVWN CSTAVCRS
18274	48642	Α	18381	1	921	
18275	48643	Α	18382	1	2115	
18276	48644	A	18383	2	149	ARRNESRRGGR*HPCRPIT*TGT VNQRPAGFRRFAGPFRYSGKPG RERL
18277	48645	A	18384	482	1655	
18278	48646	A	18385	1	776	MENACVLLMGVLSTFLVSWLG KDAMAGVGLADNFNMVIMAF FAAIDLGTTVPGAFSLCKRDRR QARVATRQSLVIMTFLPVLLAP LIHHFGEQIIDFVAGDATTEVKA LALTYLELTVLSYPAAAITLIGS GALRGAGNTKIPLLINGQVAML MQAPYYFQEAQIEAAIAAMDV APEYADIRQVESSTAVLYLFSE RFMTYGKAYGLCEWFEGSRLSI AHIPVTDNDNETQYHFFRQSCQ HC/DRCSVHRRLPDGGVVA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18279	48647	A	18386	11017	9228	FGRCAGVCQ*RGRDGQRRDEA
10277	10017	 	10300	'''	1,220	GDGRLRARL*\ALYQTITHDLR
						YRHKNLYRRKLASQRVTEHVN
						RAGLRYSYQYEKDCITITDSLA
	1					HNKVLHTQGEAGLKRVVKKK
						HADGSVTQSQFDAVGRLRAQT
						DAAGRTTEYSPDVVTGLITRITT
	1					PDGRASAFYYNHHNQLTSATG
						PDGLELRREYDELGRLIQETAP
						DGDITRYRYDNPHSDLPCATED
		İ				ATGSRKTMTWSRYGQLLSFTD
1						CSGYVTRYDHDRFGQMTAV
18280	48648	В	18387	1	1431	CSOT VICTORIDIA GQWIAV
		+	18388	575	835	RQKPERYLTAFISTTLPLRWMT
18281	48649	A	10300	575	033	LVRVMRPIVTCRRSRSILLRSIS
						HLCKWRVSTKSPVILWTILSN*
	1					RLFWKSWR*NGLQSKYQDG
10202	19650	A	18389	1207	2465	ASTAQQHQLLSPAKEETTGWRI
18282	48650	A	10309	1207	2403	FHPSDPWALGRSQGGAGLIQPP
						EVQKGSVRYRARFALLGPAMV
		1				AHRHTGRPEIRYRYDSDGRVTE
						QLNPAGLSYTYQYEKDRITITD
						-
						SLDRREVLHTQGEAGLKRVVK
						KEH\ADGSVTQSQFDAVGRLRA QTDAAGRTTEYSPDVVTGLITR
		i				1 -
						ITTPDGRASAFYYNHHNQLTSA
						TGPDGLELRREYDELGRLIQET
		1				APDGDITRYRYDNPHSDL\PCA
						TEDATGSRKTMTWSRYGQLLS FTDCSGYVTRYDHDRFGQMTA
		1		}		1
		1		}		VHREEGLSQYRAYDSRGQLIGV
		İ				KDTQGHETRFEYNIAGDLDGGI
		l				APEGG/RNG/TQYDAWGRRRTT /QGGDADDRSAGGTRARKGHT
						SFRSVGDRT\VSRHGYPGRNLR
		1				RMVRSSVPLYRLTASIAPVPQE
18283	48651	В	18390	1	2604	YRFPPHQSEEVY
18284	48652	A	18390	11	2640	
18285	48653	A	18392		549	
18286	48654	A	18393	290	389	HLCGPRTAR*ARFWHL*RAYG
10200	1000	` `				L*SGECRRAGSV
18287	48655	A	18394	1095	1268	HLCGPRTAR*ARFWHL*RAYG
10207	1.0000	1.		1.5.5		L*SGECRRAGSVAENARSACRC
						RRSAHAGRFLQTAS
18288	48656	A	18395	1056	2178	
18289	48657	A	18396	1	669	
18290	48658	A	18397	1628	1856	
18291	48659	A	18398	227	668	
18292	48660	A	18399	1	3186	
18293	48661	Α	18400	291	631	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18294	48662	A	18401	1	1437	
18295	48663	Α	18402	345	727	
18296	48664	Α	18403	870	1043	
18297	48665	Α	18404	1	2175	
18298	48666	Α	18405	1	1332	
18299	48667	Λ	18406	1	1566	
18300	48668	В	18407	1	921	
18301	48669	В	18408	1	2487	
18302	48670	В	18409	1	2919	
18303	48671	В	18410	1	2214	
18304	48672	A	18411	1714	1979	RLAWRLAGRWWAGSWRTSTII
1050.	1.0072		10			PAAACSRLRCSPPSKFCSAWCR
						WCSTRACSSCCRSTSSPMPAWF
						PSAGS*LSTARRAPPSALVLRI
			:			STOS ESTINGUIT STEVEN
18305	48673	A	18412	1201	2118	PSRQADKENRPHPGRGYPARM
10303	48073	<u>Γ</u>	10412	1201	2110	MSGLTGTITTVSTGWCTTRGH
						MKSRWSKVAIFTTRWGRRVAK
				1		RVWRRERDLTGWMSLSRKPQ\
						TWYGWDGDRLTTIQNDRTRIQ
						TIYQPGSFTPLIRRIPVLGGGDT
						MDCLRTSIRLNAASVTCAYRRI
						QISMPGSRKEVVNAREEGVEFQ
						FNFQPQYIACDEDGRLTAVGLI
						RTAMGEPGPDGRRRPPPVAGSE
						FELPADVLIMAFGFQAHAMPW
		ł				LQGSGIKLDKWGLIQTGDVGY
	1					LPTQTHLKKVFAGGDAVHGAE
		l				LVVTAMAAGRQAARDMLTLF
	1					DTK*PGTAGTATG*PQYRTTEP
		l				ASRRFISRGASRHSSEGS/SVLG
						GGDTT/MGLFADFHPPQCRQRI
		l				L\PYRRDQISMPGSRKEVVNAR
						EEGVEFQFNFQPQYIACDEDGR
		1				LTAVGLIRTAMGEPGPDGRRRI
		1				PPVAGSEFELPADVLIMAFGFQ
		1				AHAMPWLQGSGIKLDKWGLIQ
		1				TGDVGYLPTQTHLKKVFAGGE
		1				AVHGADLVVTAMAAGRQAAR
						DMLTLFDTKAS
18306	48674	В	18413	1615	1656	
18307	48675	Α	18414	1287	5056	
18308	48676	Α	18415	1	271	
18309	48677	Α	18416	1	588	
18310	48678	Α	18417	2150	2332	
18311	48679	A	18418	1	238	LEELGVIGAVLVGFSL*VLWAP
						LQLPKAGGRVKVAGIQLGEVE
						WKVS\KSCSSPDPPPGVMAAKIS
						ISQALQPPQSTR
18312	48680	В	18419	1	2721	\ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
18313	48681	Α	18420	712	1031	PVMRQRGVFRSVILIKIRSARCC
		1				QMIQPRVISHAMYSLSPFFALR
						RSTKSPKRFSMNNS*ASRSSGST
		1				SRDTSAIS*ARRSISSRTSRLLIRS
						LISLISSLLVNVV
18314	48682	Α	18421	1	1779	
18315	48683	Α	18422	693	921	
18316	48684	Α	18423	758	1183	ALTAASRSECAVIMMTGRTSCC
						DIVEKWIPARCCQMIQPRVISH
						AMYSLSPFFALRRSTKSPKRFS
						MNNS*ASRSSGSTSRDTSAIS*A
						RRSISSRTSRLLIGWRSLWYTILI
						NSIPPTNRAKSMKPEPSPNACSK
						VKRCC
18317	48685	В	18424	1	708	
18318	48686	Α	18425	1	2436	
18319	48687	Α	18426	1382	1813	
18320	48688	Α	18427	900	1028	AGYAAPGRTDRRSSPSG*GYRY
						SGCYQICRQPGRLGSRKSPV
18321	48689	A	18428	1	881	MTPSRFAKKNGSLVKRCPMGI
		ļ				DAEVTMIKPIAARPKVSHSKVR
ŀ						SYMIRLSVSFSREPITQPFSAVEI
ļ		}				QTRIRYGVSGFFLRRVDTEHHP
ŀ						DRHAARALQYHITGSNRCDAIN
						AVQFDTAGGECTNRRMAEDHG
					,	RNVAVIQMLIRLVIKQAFREVP
						SELWTNRKQACPFLDFDCGENS
						AIAANNYDWFGNMNVLTFLRD
						IGKHFSVNQMINKEAVKQRLN
						REDQGISFTEFSYNLLQGYDFA
						CLNKQYGVVLQIGGSDQWGNI
						TSGIDL/NPSSASESGVWPDRSA
			10100			DH*SRWHQIW
18322	48690	Α	18429	1	3270	AVEOUGH FURCEMOVE VEC
18323	48691	Α	18430	1	374	MKTCWQILEIESTTQIDIIRQAY
						LARLPLCHPETDPQGFKALRQA
						YEEALRLAVNPVEEADDEEKD
						AAAEHEILRAFRT/FTGFRK*SFS
				1		AFRLAEIYSAIKYLEHGGCRSIT
						LAAVCNRHRSAI

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
18324	48692	A	18431	63	1272	LRRSCNAVYCW*CVATCPS\GE EDGIVLIDQDKCRGWRMCITGC PYKKIYFNWKSGKSEKCIFCYP RIEAGQPTVCSETCVGRIRYLG VLLYDADAIERAASTENEKDLY QRQLDVFLDPNDPKVIEQAIKD GIPLSVIEAAQQSPVYKMAME WKLALPLHPEYRTLPMVWYVP PLSPIQSAADAGELGSNGILPDV ESLRIPVQYLANLLTAGDTKPV LRALKRMLAMRHYQRAETVD GKVDTRALEEVGLTEAQAQEM YRYLAIANYEDRFVVPSSHREL AREAFPEKNGCGFTFGDGCHGS DTKFNLFNSRRIDAIDHQQEMF EAIAASKNLPKEDAHALGIFLR DLTTMDPLDAQAQYSELFDRG RATSLLLFEHVHGESRDRGQA MVDLLAQYEQHGLQ
18325	48693	A	18432	2	859	WPLRRSPYSIINGSGKSAASGSV MQVAQFNSHYQYDPKFERGM YLYEHRRCFNNIIDYCNSLCYH AWLVAHKDDIERNYRGPLVQV VGWCDAIFGAEELGISRAQFDQ FLRMMQGGAQFGGGYQQQTG GGGRPGGHFFEKRRLLRMVDR LNARFTGGVLVSFKGPLLEGFG PWQGLSPWVIGPDLGLPG/CLG AWGAVSVCHCPWVCKTGSPFR WVPHRRFHPPPG*KFPENWMG FCSSRTCSGLFSHPHSPTMAFKN FELGHGTPPRGTRNRPTMSSNT YRPHRFQ
18326	48694	A	18433	479	601	CWRPLPAEKRWWYLAANWW RLAARFVFPMLCVRQ*PSSMIW TTPDADIAIGTWRSCCAVLRGR KMPVSSITMRRRCY*CWRPLPA EKRWWYLAANWWRLAARFVF PMLCVRQAAPYTK
18327	48695	A	18434	69	194	ATYENMLANPRN*KHDANRHY PPGLSCSLTVVSSRNRSARV
18328	48696	A	18435	173	298	ATYENMLANPRN*KHDANRHY PPGLSCSLTVVSSRNRSARV
18329	48697	A	18436	1	1734	

Requence	E-possible nucleotide ible nucleotide insertion) CVKGKNWPP*WRW LMLLPTLPPSVLGIR FVRQGGVWMPGD INGVVNEIWTRNHA VLASGSFFSGGLVA LGLDVLQTATRGE APQPWQQFGVTTDE
Sequence Sequence	CVKGKNWPP*WRW LMLLPTLPPSVLGIR PFVRQGGVWMPGD INGVVNEIWTRNHA VLASGSFFSGGLVA LGLDVLQTATRGE
18330	LMLLPTLPPSVLGIR PFVRQGGVWMPGD NGVVNEIWTRNHA VLASGSFFSGGLVA LGLDVLQTATRGE
LNEKLPCS LQNQLQRQ EVKKVTCK DIPLRPRFA ERNGIREPI WYKGDFFA TYARHRQQ ADLIPSPRD LNRLPNAQ 18331 48699 B 18438 1	LMLLPTLPPSVLGIR PFVRQGGVWMPGD NGVVNEIWTRNHA VLASGSFFSGGLVA LGLDVLQTATRGE
LNEKLPCSI LQNQLQRQ EVKKVTCK DIPLRPRFA ERNGIREPI WYKGDFFA TYARHRQQ ADLIPSPRD LNRLPNAQ LNRLP	LMLLPTLPPSVLGIR PFVRQGGVWMPGD NGVVNEIWTRNHA VLASGSFFSGGLVA LGLDVLQTATRGE
LQNQLQRQ EVKKVTCK DIPLRPRFA ERNGIREPI WYKGDFFA TYARHRQC ADLIPSPRE LNRLPNAQ LNRLPNAQ LNRLPNAQ LNRLPNAQ 18332 48700 A 18439 710 889 18333 48701 A 18440 1622 2615 18334 48702 A 18441 I I309 18335 48703 A 18442 I 266 PRQCGFMK PRGRCLHC YAVSCRQR VSPYTPEE M 18336 48704 A 18443 267 395 18337 48705 A 18444 I 3807 18338 48706 A 18445 I 2193 18339 48707 A 18446 1278 1846 SCRSAPRNI TTPPGYDH RDAFFAVQ	PFVRQGGVWMPGD NGVVNEIWTRNHA VLASGSFFSGGLVA LGLDVLQTATRGE
BVKKVTCK DIPLRPRFA ERNGIREPI WYKGDFFA TYARHRQC ADLIPSPRE LNRLPNAQ LNRLPNAQ LNRLPNAQ LNRLPNAQ LNRLPNAQ 18332 48700 A 18439 710 889 18333 48701 A 18440 1622 2615 18334 48702 A 18441 I I309 18335 48703 A 18442 I 266 PRQCGFMK PRGRCLHC YAVSCRQR VSPYTPEE M	NGVVNEIWTRNHA VLASGSFFSGGLVA LGLDVLQTATRGE
DIPLRPRFA ERNGIREPI WYKGDFFA TYARHRQC ADLIPSPRD LNRLPNAQ LNRLPN	VLASGSFFSGGLVA LGLDVLQTATRGE
ERNGIREPI WYKGDFFA	LGLDVLQTATRGE
WYKGDFFA TYARHRQC ADLIPSPRD LNRLPNAQ 18331 48699 B 18438 1	7
TYARHRQC ADLIPSPRE LNRLPNAQ 18331 48699 B 18438 1	
ADLIPSPRD LNRLPNAQ	` ``
LNRLPNAQ	
18332 48700 A 18439 710 889 18333 48701 A 18440 1622 2615 18334 48702 A 18441 I 1309 18335 48703 A 18442 I 266 PRQCGFMK PRGRCLHC YAVSCRQR VSPYTPEEY M 18336 48704 A 18443 267 395 18337 48705 A 18444 I 3807 18338 48706 A 18445 I 2193 18339 48707 A 18446 1278 1846 SCRSAPRNI TTPPGYDH RDAFFAVQ	
18333 48701 A 18440 1622 2615 18334 48702 A 18441 I 1309 18335 48703 A 18442 I 266 PRQCGFMK PRGRCLHC YAVSCRQR VSPYTPEEY M 18336 48704 A 18443 267 395 18337 48705 A 18444 I 3807 18338 48706 A 18445 I 2193 18339 48707 A 18446 1278 1846 SCRSAPRNI TTPPGYDH RDAFFAVQ	
18334 48702 A 18441 I 1309 18335 48703 A 18442 I 266 PRQCGFMK PRGRCLHC YAVSCRQR VSPYTPEEN M 18336 48704 A 18443 267 395 18337 48705 A 18444 I 3807 18338 48706 A 18445 I 2193 18339 48707 A 18446 1278 1846 SCRSAPRNI TTPPGYDH RDAFFAVQ	
18335 48703 A 18442 I 266 PRQCGFMK PRGRCLHC YAVSCRQR VSPYTPEEY M 18336 48704 A 18443 267 395 18337 48705 A 18444 I 3807 18338 48706 A 18445 I 2193 18339 48707 A 18446 1278 1846 SCRSAPRNI TTPPGYDH RDAFFAVQ	
PRGRCLHC YAVSCRQR VSPYTPEEY M 18336 48704 A 18443 267 395 18337 48705 A 18444 1 3807 18338 48706 A 18445 1 2193 18339 48707 A 18446 1278 1846 SCRSAPRNI TTPPGYDH RDAFFAVQ	
YAVSCRQR VSPYTPEEN M 18336 48704 A 18443 267 395	KRLENGDDYFAVN
VSPYTPEEN M 18336 48704 A 18443 267 395	CWMTVLC*RKRSD
M M M M M M M M M M	PRPPVLAPVNSISRS
18336 48704 A 18443 267 395 18337 48705 A 18444 1 3807 18338 48706 A 18445 1 2193 18339 48707 A 18446 1278 1846 SCRSAPRNI TTPPGYDH RDAFFAVQ	/KPTVRAQLEKNCN
18337 48705 A 18444 1 3807 18338 48706 A 18445 1 2193 18339 48707 A 18446 1278 1846 SCRSAPRNI TTPPGYDH RDAFFAVQ	
18338 48706 A 18445 1 2193 18339 48707 A 18446 1278 1846 SCRSAPRNI TTPPGYDH RDAFFAVQ	
18339 48707 A 18446 1278 1846 SCRSAPRNI TTPPGYDH RDAFFAVQ	
TTPPGYDH. RDAFFAVQ	
RDAFFAVQ	RHQIPGTYRSRLPRH
1) 1 1 1 1 1	NRKNPNLYISTRQK
	YHSLLYYRQTLLPN
	*FVFRL*NQNGYNS
GRHLYWRY	/HSYNAWL*LCHR
HPVYYRLY	YIFLQPVQMQMPQ
CPSPLMPEC	SFSFSFLILINA/SNI
QRESPVENI	RKVFNLRLLKIYYR
LIIFLKCGN	LLNL
18340 48708 A 18447 1 2106	
18341 48709 A 18448 881 1401	
1 1 1 1 1	VQYMASLGAPMPM
LAA/MYCG	CVCCADDDDDDDAW
	1
	AVYFLHAGYGGDW
18343 48711 A 18450 1 1960	1
! ! ! ! ! ! !	AVYFLHAGYGGDW RCGWAKYD
1 1 1 1 1 1	AVYFLHAGYGGDW RCGWAKYD AQALWHVLHY*AV
TRQLRECW	AVYFLHAGYGGDW RCGWAKYD AQALWHVLHY*AV KPIFNRGKRFRPTP
AVVARWTI	AVYFLHAGYGGDW RCGWAKYD AQALWHVLHY*AV
	AVYFLHAGYGGDW RCGWAKYD AQALWHVLHY*AV KPIFNRGKRFRPTP
AAARLSKSI	AVYFLHAGYGGDW RCGWAKYD AQALWHVLHY*AV KPIFNRGKRFRPTP IVWGSASCQLTKQ
18345 48713 A 18452 1 828	AVYFLHAGYGGDW RCGWAKYD AQALWHVLHY*AV KPIFNRGKRFRPTP IVWGSASCQLTKQ ILMRRRKGWHGRA LKQVPRQFCKPPA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	1	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18346	48714	ĪĀ	18453	<u> </u>	1327	MMQKPATVDEEESPRETHRYS
700 10	" " " " " " " " " " " " " " " " " " "					VRRRGELIVEAGGRAENTVVT
						GAGWLKVATGGIAKCTQYGN
						NGTLSVSDGAIATDIVQSEGGAI
						SLSTLATVNGRHPEGEFSVDKG
						YACGLLLENGGNLRVLEGHRA
		1				
			}			EKIILDQEGGLLVNGTTSAVVV
						DEGGELLVYPGGEASNCEINQG
						GVFMLAGKASDTLLAGGTMN
1						NLGGEDSDTIVENGSIYRLGTD
1						GLQLYSSGKTQNLSVNVGGRA
						EVHAGEVTGDIHVEITDASRQT
						LCNALKLQPDEGGSSRVIPTCK
						ASRLRIDANFKRFVDEEVLPGT
						GLDAAAFWRNFDEIVHDLAPE
						NRQLLAERDRIQAALDEWHRS
		İ				NPGPVKDKAAYKSFLRELGYL
						VPQPERVTVETTGIDSEITSQAG
						PQLVVPAMNARYALNAANAR
			}			WGSLYDALYGSDIIP\QEGAMV
			·			SGYDPQREAMECP
18347	48715	В	18454	71	4995	
18348	48716	Α	18455	1	3572	MARKRKSRNNSKIGHGAISRIG
		İ				RPNNPFEPCRNRYAQKYLTLAL
		ŀ				MGGAAFFVLKGCSDSSDVDND
		ĺ				GDGTFYATVQDCIDDGNNADI
		ĺ				CARGWNNAKTAFYADVPKNM
ŀ						TQQNCQSKYENCYYDNVEQS
						WIPVVSGFLLSRVIRKDRDEPFV
						YNSGGSSFASRPVWVPSSIVIMS
						YGLASYVIAVAFLERDRIQAAL
						DEWHRSNPGPVKDKAAYKSFL
						RELGYLVPQPERVTVETTGIDS
		İ				EITSQAGPQLVVPAMNAR
18349	48717	A	18456	485	1999	ETTSQACT QUV VI AMITA IK
18350	48718	Α	18457	44	270	
18351	48719	Α	18458	1	2223	
18352	48720	Α	18459	290	756	
18353	48721	Α	18460	8	314	
18354	48722	A	18461	2	686	
18355	48723	Α	18462	1	732	
18356	48724	Α	18463	1	780	
18357	48725	Α	18464	101	435	
18358	48726	A	18465	805	993	
18359	48727	A	18466	452	2180	
18360	48728	A	18467	691	1443	DDD DOD L KVOKEN KEEK FOROM
18361	48729	Α	18468	1	320	DPRDCRLKVGKEMFTLFGPQF
•						VRELQQRGFDIFLDLKFHDIPNT
-						AAHAVAAAADLGVWMVNVH
						ASGGARMMTAAREALVPFRPD
L			<u> </u>		<u> </u>	APLLIAVPV*RSIKASDLVVPL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence		in USSN 09/540,217	location of first codon for peptide sequence		*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
18362	48730	A	18469	632	796	
18363	48731	A	18470	129	425	TDIITRQNGTGANIDAIEIIATKA STGLFVYQPRFARCPRVMSSGY HTATRGSSGTRTVRARVCHTA AGVYGSSKQTQF*NRFCQRKCR
		<u> </u>				RYPDGCRRA
18364	48732	Α	18471	1	1062	
18365	48733	A	18472	1	996	
18366	48734	A	18473		1004	MLRVYHSNRLDVLEALMEFIV ERERLDDPFEPEMILVQSTGMA QWLQMTLSQKFGIAANIDFPLP ASFIWDMFVRVLPEIPKESAFN KQSMSWKLMTLLPQLLEREDF TLLRHYLTDDSDKRKLFQLSSK AADLFDHVSGASVSSKVEQLR AQLNERILVLDGGMGTMIQSY RLNEADFRGERFADWPCDLKG NNDLLVLSKPEVIAAIHNAYFE AGADIIETNTFNSTTIAMADYQ MESLSAEINFAAAKLARACADE WTARTPEKPRY\VAGVLGPTNR TASISPDVNDPAFRNITFDGLVA AYRESTKALVAGIVLMPAIPAL WEAEPGGCA
18367	48735	A	18474	606	846	
18368	48737	A	18475	3042	1639	GPHPPAPGGPA\CHYDVHQPYL RQRTEDRRTRHYLAQSFAQYSL RQWAFHFLLDNITPDIQQIVIAH TRRTGGFTITASEAAIEMLLCFK RDFVALQHLFNQIDAPARAVQF IAQKLIGWACCEAARKNQLPH NTTKRSLLMSILYEERLDGALP DVDRTSVLMALREHVPGLEILP TDEEIIPYQCDGLSAYRTRPLLV VLPKQMEQVTAILAVCHRLRV PVMVFPGLRLSREAFLTTAEFG TQELRRPAQLRLQVYRLLSTAG RHHPAQNSPIRVLEVDIDLLGG TQELWGWKETH ICSRCKSTRTSPTRCRKAIKVIL
18309	48/3/	A	104/0	009	1039	ICSRCRSTRTSPTRCRRAIKVIL VCFVIRELLVVAPELTIIGLPEG NWITVSKGMLAKSPVDSNSTQ LKEAEEERLKAAQYGLQLVES QNELQNQLDKCRNEMMTMTE MQLRNKKRRRELGNLIKRFRSG AESLPDAVVLTTEEGGIFWCNG LAQQILGLRWPEDNGQNILNLL RYP/EVYAISENA*FFSPAQSGA QHRAASGNSRHALYPQTVADG GA*CHANASTGRGAA*LFCQRE P*VTYAIDRVTGLPGDDE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence		*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
18370	48738	A	18477	388	2356	
18371	48739	A	18478	137	318	GGQLSSWAMDFCHL/PAFWNS
10371	1.0735		10770	1.57		MTWHLQRLWGACGYFWHAQ
						WERLLTTFEGNEWILIFIGGIL
18372	48740	A	18479	923	1149	
18373	48741	Α	18480	1	373	
18374	48742	A	18481	168	360	
18375	48743	Α	18482	1	147	
18376	48744	Α	18483	161	440	
18377	48745	Α	18484	277	2172	
18378	48746	A	18485	164	406	LFRREGSYDRYLLMDDYCRRK DDSYFDRYRDSFDGRGPPGPES QSRAKGKVAETVYSYY**NTL QKREETNNQYPHLKKG
18379	48747	A	18486	405	999	
18380	48748	Α	18487	64	288	
18381	48749	A	18488	159	550	CAVCHISFQDTPVLVSSNVTMQ FGSKPLFENISVKFGGGNRYGL\ IGANGSGKSTFMKILGGDLEPT LGNGSLDPNERIGKLRQ\DQFAF EEFTVLDTVIMGHKELWEVKQ ERDRIYALPEMSEEDGYKVAD
18382	48750	Α	18489	3	481	
18383	48751	A	18490	2	233	
18384	4875 2	A	18491	2714	3026	SVPTIIFFFLPVLRFWRPGIQQA KKCHLFATGTQSRRNTLRQRT AKGIPQQRMTAKREEISIGILHV TPQQRRECRGDNRFTARFRRTP WRLLGHCVSAAVTGVLPAVAG SPLAYSDTDEFYPVAGGTMSQ HLPLVAAQPGIWMAEKLSELPS AWSVAHYVELTGEVDSPLLAR AVVAGLAQADTLRMRFTEDNG EVWQWVDDALTFELPEIIDLRT NIDPHGTAQALMQADLPQDLR VDSGEPLVFHQTTLGFI RLIGLRMVREFVGLIRRASVAS
						GSWHCCRMRRKRLIRPKPSILQ SMLWSPSTRRMFFTLVPIFSIEE LPFTFRSLMMVTLSPSASRLPLA SLTISSSLSCAGSASFHS*AHSG QMYMPLSS
18386	48754	Α	18493	1	888	
18387	48755	A	18494	1778	2025	
18388	48756	A	18495	1	1425	DVGD IDDI DDIDDI/*EMDI MD ** C
18389	48757	A	18496	3	96	FVGRIRRLRRIRRY*FTPLTD*LS CAHQVP
18390	48758	Α	18497	420	955	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18391	48759	A	18498	533	1006	RPSIFACVSP*RINHSRIMPQTN
	10,05					AKNAKPNPAQVMRGVISRRRR
				1		AMLFSNSAITRTPTTSFTMAITS
						ASSSPPAVKVSICPHGINTPTNIC
						NTTSCFRLIPHFMTPKPLPLYSE
						SGPSCSSCSASARSNGSLPISIKV
			1			ANAQAQSASTSGMLWLSGQ
18392	48760	A	18499	3	418	EWHYLSADRRQCAFQGVSRKT
10372	10700	` `	10155	ľ		ATVCHHPVDYYAGSLYRLYFT
						DRLRARLAGHAAES/GTPASHA
						VFQLVLE*L*SPLFSPVSTSGGR
						TANSTVLIMKSCMSGAVERQPT
						NWQAILMFLIFVVFTLGITYWA
						SNAYVLAG
18393	48761	C	18500	63	500	5
18394	48762	Α	18501	1	1107	
18395	48763	Α	18502	1	153	
18396	48764	Α	18503	88	510	
18397	48765	Α	18504	204	654	
18398	48766	Α	18505	241	771	
18399	48767	Α	18506	1655	1909	
18400	48768	Α	18507	2	48	
18401	48769	Α	18508	1	585	
18402	48770	Α	18509	1	723	LQPPGRRWAAPAV\SGLSRQV\
						RCFSTSVVRPFA\KLVGPS\VQV
						SVIEGRYATAL\YYAASRQNKL
	Ì					EQVEKG\LLRVAQILEGTPKWA
						ASVLESPMLKAFPLKVKKPKM
						TITAKVGGSLPLTYQTLINFALL
1		1				ENGSD*GNYPRESFSGLFLTMD
1						GVFHSRERVP\CTV\TSGIFL*EG
		1				RPQFSEFKNCSSRGFLRSRAKFL
		1				KLGAK\TDPSI\LGGMIVRIGE\K
						YVDMSVKTK\IQKLG\RAMREI
18403	48771	Α	18510	400	894	
18404	48772	Α	18511	1	1023	
18405	48773	Α	18512	73	424	RPGMWSTRSPNSTAWPLSLEPD
		1				PGMASACTTMHTTTIAEPDPG
1						MSGWPDGRMETSSPTIMDIVVI
		l				*CAIAAEAIVLVSLLFVMLRYM
		l				YRHMGTYHTNEAKGTGVADS
						ADACPAGDPA
18406	48774	A	18513	141	563	RPGMWSTRSPNSTAWPLSLEPD
		1				PGMASASTTMHTTT\IA\EPDPG
		ł				VS\GLP\DGRMETPTPHP*LTMV
		1				VIAGV\IACLWPFVLSLPSSFVM
						VALTLLTGHKGARTTPNEAKG\
						TEFAES\ADAALQG\DPAL\QDA
	1					GDSSRK\EYFI

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
18407	48775	A	18514	182	935	QAFELHCPCRPKFSTLPSPAWPP RHTDLRKRGRRTAQ*YAGNES HPPSLPRYLRRSRH/SRMQAPSS TCSYSHTSLQCSPAPQNH/AQHP *PALEELACG/SPSVSSPYLVLSS CQEQPHHCCPPSTPRPSW/CPSS GDAICYSPGQ/CPRSRGPLP\EDS SDSPPAEQVLPPSSGSHNTLYLR CKRFSAFILNCEPPSK\C*KPGHR CRS*AGILTLLPPG/ERGP/GDGP CSTGRQSASKTPPPPSHPHTGHS LWSEEK
18408	48776	A	18515	48	676	KARLTWQQTREESLCRETPPYK TIRSRETYSLPREQHGKDLPP*F NYLPPSPSHNTICPG*QDQGRFT Q\PLYLRCKRFSAFILNCEPPSK\ C*KPGHRCRS*AGILTLLPPGGE AQHFKVKHPSIPL/REGGEVGIK PKKTSPGTELLMQP/CHQPSPRT CQPAPRKSNRP/LGCLSPLMKHS TLGYLLWLQKAIWLKGFGTSG W*M*GRFHL
18409	48777	A	18516	86	92	TANAALHQPSPRTCQPAPRKSN RPQAA*ALNC
18410	48778	В	18517	1	2031	
18411	48779	A	18518	98	367	DLFLYLSWQHCHPYSHCHPNQ KPGRFLMKLSS\ETVTIELKNGT TRSHGT\IHQVWDVQ\MNTH\L KAVKMTLKNREPVQL\ETLSIR GNNIR\YFILPD/SVYPLDTLLVD VDPKVKSKKREAVAGRGRRRG RG/TRDVARGR*RGNNIR
18412	48780	Α	18519	1	310	
18413	48781	A	18520	37	136	RRRPSSSWALPPSRGPSST*AER GAQHQQPVPA
18414	48782	A	18521	3	305	FFTLCLQWGLFSCESPICSGVGR YWA*VCLCFLQIEAIELPMDPK LNKRRGFVFITFKEEEPVKKVL EKKFHTVSGSKVRCSQLCLASC AAGELALRGMG
18415	48783	В	18522	249	367	
18416	48784	Α	18523	1	1008	
18417	48785	В	18524	17	835	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
İ	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18418	48786	Α	18525	2	1188	SLIKLNSPPPRGGGTARYGAWL
10410	140700	l'`	10323	[1100	LVGGFPCGGGQGGDTVGAAP
						WADWGEASAAASFDVLAEAC
						ARLSSLRVGDEAGSRGVIWRAR
						SPPRLGLALSEAG\EE\QPMETT
						GASENGHEAVPE\ASR\GRGCW
						TPLRRGLEARPRRPRAGIRTAPR
						D/QRINASKNEEDAGKMFVGGL
						S\WDTSKKDLKDYFTKFGEV\V
						DCTIKMDPNT\GRSRGFGFILFK
1						DASSV*K\VLDQKEHRLDGRVI
						DP\KKAMA\MKKAPVKKIFVGG
						LNPESPTEEKIRESFGEF\GEIEAI
						ELPMDPKLNKRRGFVFITFKEE
						EPVKKVLEKKFHTVSG\SKCEIK
						VAQPKEVYQQQQYGSGGRGN
						RNRGNRGQ/GGGGGGG*GQGS
			1			TNYG\KTQ\RRGGHQ\NNYKPIL
						EAGRQERPT
18419	48787	Α	18526	3	784	
18420	48788	Α	18527	3	1990	PAMNGLSLSELCCLFCCPPCPG
		1				RIAAKLAFLPPEATYSLVPEPEP
		l				GPGGAGAAPLGTLRASSGAPG
						RWKLHLTERADFQYSQRE\RST
						IEVFPTKSARGNRVSCMYVRCV
		l				PGARYT\VLFSHGNAVDLGQMS
						SFYIGLGSRLHCNIFTYDSSGYG
						ASSGRPSERNLYADIDATWQAL
						RTR*GRPLVGRVRARWRPRLTL
						LRRRQVRHQPGQHPSCTGRSIG
						HGAHRGTWPRATSVPRWCCTR
						RSTSGMRVAFPDTQERPTASTP
						FPYIRERCPRLTSPRAHHPTGTK
						DEVIDFLEAGWRSTRRCHQGG
		İ		ŀ	ļ	GAAVGGGGPGHNDIELYSQYL
						ERLASLSFSQELPSQRA
18421	48789	Α	18528	82	406	

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
18422	48790	A	18529		1474	ARASGSGSSSGQGKMGQS\QRG GHGSGGKKDDKDKKKKYEPP VPTRVGKKKKKKTKGPDAASK\L PLVTPHTQCRLKLLKLERIKDY LLME\EEFIRNQEQMKPLEEKQ EEE\RSKVDDLKGGPRLSVGNL GKRFI\DDNHA\IVSTP\VGSEHY\ VSIL\SFVDKDLLEPGCSVLLNH KVHAVIGVLMGDTDPLVTV\M KVEKAPQETYADIGG\LDNQI\Q EIKESVELPLTHPEYYEEMGIKP PKGV\ILLVPPG\TGKTFLPKAV ANQT\SATFLRGGGSELIQKNLG DGPKT/LSRELFR\VSEELAPSLR FIDRILPPLGTKRY\DSNSGGE\R EFQRTM\LELLN\QL\DGFDS\RG D\VKVIMPTNR\IETLDPALIRPR PAFDRKYEFP/LCPDEK\TKKRIF QISHKS/RMTLADVGTPWTTLI MA\KDDLSGA*PSKAI\CT\EAGL MAL/RENRRMKVPN\EDFKKS\ KENVLYKKQEGTPEGVVSLMN
18423	48791	A	18530	2	274	PWLVIRKMVGRFLNP WQDLICRMAASVKEQSTKPIPL PQSTPG/ESHSLSTSGKSEVRDL FVAERQFAKEQHTDGTLKEVG EDYQIAIPDSHLPVSEERWALD ALRN
18424	48792	Α	18531	3	844	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
18425	48793	A	18532	261	2814	VRFRNLIRIYCNIKICAVSHCPP
						KVFKLNSLFFLHRFDSVAFGES
						QSEDEQFENDLETDPPNWQQL
						VSREVLLGLKPCEIKRQEVINEL
						FYTERAHVRTLKVLDQVFYQR
						VSREGILSPSELRKIFSNLEDILQ
	1					LHIGLNEQMKAVRKRNETSVID
						QIGEDLLTWFSGPGEEKLKHAA
						ATFCSNQPFALEMIKSRQKKDS
						RFQTFVQDAESNPLCRRLQLKD
						IIPTQMQRLTKYPLLLDNIAKYT
						EWPTEREKVKKAADHCRQILN
						FVNQAVKEAENKQRL/EEDYQ
						RRLDTSSL\KLSEYPNVE\RLRN
						MDLTKRKMIHEGPLVWKVNR
						DKTIDLYTLLLEDILVLLQKQD
						DRLVLRCHSKILASTADSKHTF
						SPVIKLSTVLVRQVATDNKALF
						VISMSDNGAQIYELVAQTVSEK
						TVWQDLICRMAASVKEQSTKPI
						PLPQSTPGEGDNDEEDPSKLKE
						EQHGISVTGLQSPDRDLGLEST
		ĺ				LISSKPQSHSLSTSGKSEVRDLF
						VAERQFAKEQHTDGTLKEVGE
						DYQIAIPDSHLPVSKE\RRALDA
						LRNLGLLKQLLVHQLGLTEKSA
						LENWQHFPRYRTAFQGPQTDS
		1	ĺ			VIQNSENIKAYHSGEGHMPFRT
						GTGDIATCYSPRTSTESFAPRDS
						VGLAPQDSQASNILVMDHMIM
						TPEMPTMEPEGGLDDSGEHFFD
						AREAHSDENPSEGDGAVNKEE
18426	48794	Α	18533	90	383	ILQLHSSGPCLLWL*LSFRSLST
						TAV\CQCCRPAADFHPSGSSRVS
						AVLLIQ/HRRPLPLPIGLKACYFS
		1				ALSLQTSLTCLGLVVLLDFPKE
	L	<u>L</u> _				AASLDKV
18427	48795	Α	18534	1	456	
18428	48796	Α	18535	830	1092	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18429	48797	A	18536	391	1824	L GSGADASGFISDAGLIFKKASSK
						GPKRLEKFSDERAAYFRCYHKL
						FLLFSSLCYGTYHVVVLPTLKP
						REDEDWISLCLRHCLMQKDTH
						SMLGECGERREGKKQKTGALG
		İ				SSLSLEDAKHLLPFIPALPFSWN
						ALPQDLCNVFKLLWDYRIRVD
			ŀ			PSPIAGTLIRRDTDPQGESHVKT
						GRHWSDTIKSQRCQGLRQPPDR
						PIPILLMVKGQHVWPSQLDKAQ
						IQEVTELNNVKNVARLPKSTKK
						HAIGIYFNDDTSKTFACESDLEA
						DEWCKVLQMECVGTRINDISL
		l				G\EPDLL\ATGVE\REQSE\RFNV
						YLMTSP*LRMLHG\ECALQV\T
						Y\EYICLWGRPRTPQEFKLISW\P
						LSALRRY\GRGTPWFNYQAGR
						M\CQTGEGLFIFQTRDGEAIYQK
			[VHS\AALAIAEQHERLLQSVKN
]			SMLQMKMSERAASLSTMVPLP
		ŀ	1			RSAYWQHITRQHSTGQLYRLQ
		ŀ				DVSSPLKLHRTETFPSLQDS
18430	48798	C	18537	22	620	
18431	48799	Α	18538	48	413	EDSKSKTISHPWRDCGDYCHH
				1		QGLERCRGGDSHHTPVQLS*LA
		ŀ				CAED\NKSWRMT\GIIVSLTKC*
					1	LQMQLLYQMWFHCLSKLTHLL
						GISTLQLCVTILFREILITFPLHKI
						SHWSITLMTLR
18432	48800	A	18539	23	124	QWVIISLTKW*LQLQLLYQMW
						FHCLSKLTRLLP
18433	48801	Α	18540	1	1860	
18434	48802	Α	18541	209	3816	QGRPTFRFRKYREHHKDTPREE
						QLQDT*SSDSPKLK*RKKC*GQ
						PERKVKLPTKGSPSD*KRISRQ/
						KTLQARRQSWFFEKINKIDRPQ
	:					ARLIKKKREKNQIDTIKNDKGD
						ITTDPTEIQITIREYYKHLYANK
						LENLEEMDKFLDTYTLPRLNQE
		l	•			EVESVNRPITGSEIEAITNSLPTK
						KSPGPDGFTAEFYQRYKEELVP
						FLLKLFQPIEKEGILPNSFYEASII
		ļ				LIPKPGRDTTKKGNFRPISLMNI DAKIL
18435	48803	Α	18542	1	1521	
18436	48804	В	18543	1	2265	
18437	48805	В	18544	1	1716	
18438	48806	Α	18545	1	1704	
18439	48807	A	18546	1	3618	
18440	48808	Α	18547	1	3087	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence		in USSN 09/540,217	location of first codon for peptide sequence		*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
10441	40000		10540	1077	350	 SPVEKELKLWKNRHKLLSC*W
18441	48809	Α	18548	277	330	LTC
18442	48810	A	18549	197	422	IVNKVVLPQGGPASPSLKGFWV
101.2	10010		103 17			YKLAQVWKLIEGP*FPIFIIQISL
						LSI*PRSFLFV*IK*ECSRLPISFTS
		ļ				SGR
18443	48811	A	18550	1	3171	MVKGSIQQEELTILNIYAPNTG
		ł				APRFIKQVLSDLQRDLDSHTLI
						MGDFNTPLSTLDRSTRQKVNK
						DTQELNSALHQADLIDIYRTLH
						PKSTEYTFFSAPHHTYSKIDHIL
						GSKALLSKCKRIEIITNYLSDHS
						AIKLELRIKNLTQSRSTTWKLN
						NLLLNDYWVHNEMKTEIKMFF
						ETNENKDTTYQNLWDAFKAVC
						RGKFIALNAYKRKEERSKIDTL TSQLKELEKQEQRHSKPSRRQE
						ITKMRAELKEIETQ
18444	48812	A	18551	209	2432	QGRPTFRFRKYREHHKDTPREE
10444	70012	1	10331	200	2-132	QLQDT*SSDSPKLK*RKKC*GQ
						PERKVKLPTKGSPSD*KRISRQ/
						KTLQARRQSWFFEKINKIDRPQ
						ARLIKKKREKNQIDTIKNDKGD
				:		ITTDPTEIQITIREYYKHLYANK
				1		LENLEEMDKFLDTYTLPRLNQE
						EVESVNRPITGSEIEAITNSLPTK
		:				KSPGPDGFTAEFYQRYKEELVP
						FLLKLFQPIEKEGILPNSFYEASII
						LIPKPGRDTTKKGNFRPISLMNI
		ŧ				DAKILNKILANQIQQHIKKLIHH
						DQVGFIPGMQG\WLEVLARA/IR
						QEKEIKGIQLGKEEVKLSLFAD
		ŀ				DMIVYLENPTVSAQNLLKLISN FSKVSGYKINVQKSQAFLYTNN
						RQTESQIMSELPFTIASKRIKYL
						GIQLTRDVKDLFKENYKPLLNE
						IKEDTNKWKNIPCSWIGRINIVK
						MAILPKVIYTFNAIPIKLPMTFFT
						ELEKTTLKFIWKQKRSHIAKSIL
						SQKNKAGDITLPDFKLHYKATV
	ĺ					TKTAWYWYQNRDIDQWNTTE
						PSEIMLHIYNHLIFDKPDKNKQ
						WGNDSLFNKWCWENWLAICR
						KLKLDPFLTSYTKINSRWIKDL
						NVRPKTIKTLEENLGNTIQDIG
						MGKDFMSKTPKAVATKAKIDK
						WDVIKLKSFCTAKETTIRVNRQ
						PTEWEKIFATYSSDKGLISRIYN
]						ELKQIYKKKTNNPIKKWAKDM
	<u> </u>					NRHFSKEDIYAAKRHMKKCSSS
18445	48813	Α	18552]1	3255	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	1	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18446	48814	A	18553		3139	
18447	48815	$\frac{\Lambda}{\Lambda}$	18554	1	3210	MVKGSIQQEELTILNIYAPNTG
10447	40013	^	10004	1	3210	1
						ALRFIKQVLRDLQRDLDSHTIIM
						GDFHTPLSTLDRSTRQKVNKDI
						QELNSALHQEDLIDIYRTLHPKS
		1			:	TEYTFFSAPHHTYSKIDHIVGSK
	1					ALLSKCKRTEIITNCLSDHSAIK
İ		1				LELRIKNLTQNRSTTWKLNNLL
						LNDYWVHNEMKAEIKMFFETN
						ENKDTTYQNLWDTFKAVCRGK
						FIALNAHKRKQERSKIDTLTSQL
						KELEKQEQTHSKASRRQEITKIR
						AELKEIETQ
18448	48816	Α	18555	1	3345	
18449	48817	В	18556	1	3127	
18450	48818	A	18557	1	3325	
18451	48819	Α	18558	1	3145	
18452	48820	Α	18559	1	548	
18453	48821	A	18560	1	3229	
18454	48822	Α	18561	1	3352	
18455	48823	Α	18562	1	1965	MGDFNTPLSTSDRSTRQKVNK
				İ		DTQEMNSALHQADLIDIYRTRH
						PKSTEYTFLSAPHHTYSKTDHIL
						GSKALLSKCKRTEMITNYLSDH
						SAIKLELRIKKLTQNCSTTWKL
						NNLLLNDYWVHNEMKAERKM
						FFETIENKDTTYQNLWDAFKAV
						CRGKFIALNAHKRKQERSKIDT
						LTSQLKELEKQEKTHSKTIRRQ
						EITKIRAELKEIETQKTLQKINES
						RSWFFERINKIDRTLARLIKKKR
						EKNQIDRIKNDKGDITTDPTQIQ
						TTIREYYKHLYANKPENLEEMD
		1				KFLDTYTLPRLNQEEVESLNRPI
						SGSEIVAIINSLPIKKSPGPDGFT
						AEFYQSYKKELVPLLLKLFQSIE
						KEGILPNSFYEASIILLPKRGRDT
					1	TKKENFRPISLMNIDAKILNKIL
		ĺ				AKRIQQHIKKLIHHDQVGFIPG
						MQ\GIKYLGIQLTRDMKDLLKE
						NYKQLLNEIKEDTNKWKNIPCS
						WVGRINIVKMAILP/KELEKTTL
				1		KFIWNQKRACIAKSILSQKKKA
				1		GGITLPDFKLYYKATVTKTARL
				1		YTKHGTSICCGSFGELLPMQKA
				1		_
				1		KRELAHHMAKAGARVKGQLG
				1		HVTQHQLGSIEWFNKYSLSTYY
						TVDTKPQFSHPTKDNKTGLTCR
				1		NRTTVGVDLQAKGNLVVKHLS
		<u></u>		<u> </u>	<u> </u>	SLQHCYEFICHI

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
18456	48824	A	18563	1	3099	MGELITPLSTLDRSTRQKVNKD TQELNSALHQGDLIDIYRTLHP KSTEYTFFSAPHHTYSKIDHILG SKALLSKCKRTEIITNYLSDHSA IKLELRIKNLTQNRSTTWKLNN LLLNDYWIHNEMKAEIKMFFET
						NENKDTTYQNLWDAFKAVCR GKFIALNAHKRKQERSKIDTLT SQLKELEKQEQTHSKASRRQEI TKIRAELKEIETQKTLQKINESR SWFFERINKIDRPLARLIKKKRE KNQIDTIKNDK
18457	48825	A	18564		3253	MGDFNTPLSTLDRSTRQKVNK DTQELNSALHQGDLIDIYRTLH PKSTEYTFFSAPHHTYSKIDHIV GSKALLSKWKRTEIITNYLSDH SAIKLELRIKNLTQSRSTTWKLN NLLLNDYWVHNEMKAEIKMFF ETNENKDTTYQNLWDAFKAVC RGKFIALNAHKRKQERSKIDTL TSQLKELEKQEQTHSKASRRQE ITKIRAELKEIETQKTLQKINESR SWFFERINKIDRPLARLIKKKRE KNQIDTIKNDK
18458	48826	A	18565	1	3095	
18459	48827	A	18566	2	2678	

SEQ ID NO:	SEQ ID NO: of peptide sequence	•	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	B contract of the contract of	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
18460	48828	A	18567		2478	MKAEIKMLFETKENKDTTYQN LWDALKAVCRGKFIALNAHKR KQERSKIDTLTSQLKELEKQEQ TLSKASRRQEITKIRAELKEIET QKTLQKINESRSWFFERINKIDR PLARLIKKKREKNQIDAIKNDK GDITTDPTEIQTTIREYYKHLYA NKLENLEEMDKFLDTYTLPRLN QEEVESLNRPITGAEIVAIINSLP TKKSPGPDGFTAEFYQRYKEEL VPFLLKLFQSIEKEGILPNSFYE ASIILIPKPGRDTTKKENFRPISL MNIDAKILNKILANRIQQHIKKL IHHDQVGFIPGMQGWFNIRKSI NVIQHINRAKDKNHMIISIDAEK AFDKIQQPFMLKTLKKL\EIKYL GIQLTKDVKDFFKENYKPLLKE IKEDTNKWKNIPCSWVGRINIV KMAILP/KELEKTTLKFIWNQK RACIAKAILSQKNKAGGITLPDF KLYYKATVTKTAWYWYQNRD IDQWNRTETSEITPHIYNYPIFD KPEKNKQWGKDSLFNKWCWE NWLALCRKLKLDPFLTPYTKIN SRWIKDLNVRPKTIKTLEENLGI TIRDIGMGKDFMSKTSTAMAT KAKIDKWDLIKLKSFCTAKETT IRVNRQPTKWEKIFATYSSDKG LISRIYNELKQIYKKKTNNPIKK WAKDMNRHFSKEDIYAAKKH MKKCSSSLAIREMQIKTTMRYH
19461	48820	D	19569	1	2402	LTPVRMAIIKKSGNNRTPTRPIQ
18462	48829 48830	A	18568	2	3403 2976	APRFIKQVLSDLQRDLDSHTIIM GDFNTPLSTLDRSTRQKVHKDT QELNSALHQADLIDIYRTLHPK STEYTCFSAPHHTYSKIDHIVGS KALLSKCKRSEIITNCLSDHSAI KLELRIKKLTQNRSTTWKLNNL LLNDYWVHNEMKAEIRMFFET NENKDTTYQNLWDTFKAVCRG KFIALNAHKRKQERSKIDTLTS QLKELEKQEQTHSKASRRQEIT KIRAELKEIETQKTL\QKINE\SR SWFF\ERINK
18463	48831	A	18570	472	829	AIASESASPKAWQLPCGAAQQF AALPGLGNYKFPLTLITGYGSN KSPP*GYLVFQTYSSLPLLRLPP GHFGFLLPLSQ\RLRRELQCWL G*LTQTIKMKSVYHSTTEVRKS MHGIQEIH

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
	of peptide	hod	in USSN	location of first	codon for last amino acid	1
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
18464	48832	В	18571	1	2293	
18465	48833	Α	18572	1	3810	
18466	48834	В	18573	1	4107	
18467	48835	В	18574	87	3161	
18468	48836	Α	18575	1	3192	
18469	48837	A	18576	1	1983	
18470	48838	Α	18577	1	3156	
18471	48839	В	18578	1	3471	
18472	48840	В	18579	1	2337	
18473	48841	Α	18580	1	3852	
18474	48842	Α	18581	1	3057	
	48843	A	18582	1202	2832	SGIYMIGFEQVLKAQPPLSSPNG PMNKVATVVGMEVMHGLSNM DFHSRRLTWLRPLPGFTGPG/Y QGVKVEVAPLTITPSDPLAKFL LLISATSTSLEVTVPEGEMLPPR DTTKIATFGTQTGFLALQLADG LLWDLVIIPGKGKPSRDLVESPS PYSTYEGIDGWPDEPAPTATKP PVMPAPALPPDTRSGSKAPTVP TPYPQMEHHQVQLASNNSNTE ALGHLSPQSSWVQTPGQNSGP AIPNHLGKDMISPPQMAPAGVK WESQKY
18477	48845	A	18584		3702	MQKKHDKIRNSFMIKTLKKLRT EGIYLNTIKATYKKPITNIILNGE NQKAFALRSGTRFWILALCQM SRLRKLTNYLSDHSAIKLELRIK NPTQSRPTTWKLNNLLNDYW VHNEMKAEIKMFFETNENKDT TYQNLWDAFKAVCRGKLIALN AHKRKQERSKIDTLTSQLKELE KQEQTHSKASRRQEITKIRAEL KETETQKTLQKINESRSWFFERI NKIDRPLARLIKKKREKNQIDTI KNDKGDITTDPT
18478	48846	A	18585	1	3139	
	48847	A	18586	2	2918	
	48848	В	18587	1	3242	
	48849	A	18588	1	3640	
	48850	A	18589	371	522	
	48851	В	18590	1	3918	
		↓				
18484	48852	Α	18591	1	3057	

hod hod hod hod hod A	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
		sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
53 A	18592			
53 A	18592	1		
	10372	11	3233	MGKKQSRKTANSKKQSTSPPP
	E	'	3233	KEHSSSPATEQSWTENDFDELR
į				EEGFRRSNYSELREDIQTKGKE
Ł				VENFEKNLEECITTITNTEKCLK
				ELMELKAKAQELREECRSLRSR
				CNQLEERVSAMEDEMNEMKRE
				GKFREKRIKRNEQSLQEIWDYV
				KRPNLRLTGVPESDGENGTKLE
				NTLQDIIQENFPNLARQANIQIQ
				EIQRMPQRYSSRRATPRHIIVRF
				TKVEMKEKMLRAAREKDRSTR
				QKVNKNTQELNSA
54 A	18593	965	4165	TWKGTTSTSRCKIMPKYRSTRQ
		ŀ		KVNKDTQELNSALHQADLIDIY
				RTLHTKSTEY/TFFSAPHHTYSK
				IDHIVGSKALLSKCKRTEIITNY
				LSDHSAIKLELRIKNLNQSRSTT
				WKLNNLLLNDYWVHNEMKAE
				IKMFFETNENKDTTYQNLWDA
1				FKAVCRGKFIALNAHKRKQERS
ł				KIDTLTSQLKELEKQEQTHSKA
:				SRRQEITKIRAELKEIETQKTLQ
				KINESRSWFFERINKIDRPLARLI
				KKKREENQID
55 A	18594	1	5093	MEGEMNEMKREGKFREKRIKR
İ				NEQSLQEIWDYVKRPNLRLIGV
				PESDGENGTKLENTLQDIIQENF
				PNLARQANVQIQEIQRTPQRYS
				SRRATPRHIIVRFTKVEMKEKM
				LRAAREKDRSTRQKVNKDTQE
				LNSALHQADLIDIYRTLHLKSTE
				YTFFSAPHHTYSKIDHILGSKAL
				LSKCKRTEIITNYLSDHSAIKLE
				LRIKNLTQNRSTTWKLNNLLLN
				DYWVHNKMKAEIKMFFETNEN
	1			KDTTYQNLWDTF
56 A	18595	1	131	
57 A	18596	5382	9269	RAKSPANIIMTGSNSHITILTLN
				VNGLNSPIKRHRLASWIKSQDP
				SVCCIQETHLMCRDTHRLKIKG
				WRKIYQANGKQKK\AGVAILVS
				DKTDFKPTKIKRDKEGHCIKVK
				GSIQQEELTILNIYAPNTGAPRFI
				KQVLSDLQRDLDSHTLIMGDFN
	1			TPLSTLDRSTRQKVNTDTQELN
				SALHQADLIDIYRTLHPKSTEYT
				FFSAPHHTYSKIDHIVGSKALLS
				KCKRTEIITNHLSDHSAIKLELRI
				KNLTQS
	18597	L	l	
	56 A	55 A 18594 56 A 18595	55 A 18594 I	55 A 18594 I 5093 56 A 18595 1 131

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18491	48859	Α	18598	142	876	RKKGILPPKESLKELEEEAEEEQ
						RILQQSVVKTYEDMTLEELEDH
						EDEFNEEDECAIEMCRQQRLAE
	,				:	WKATKLKNKFGEVLEISGKDY
		l				VQEVTKA/ERRALWVI\LHLYK
						QGIPLCAL\VNQHLSWTLPGSFP
						DVKFIKAISTTCIPNYTDRNLP\T
						ILVYPGRRYQGLSLLVLWCLAG
		1				MNPDQEMSWEWKLSESGAIMT
						DLEENPKKPIEDVLLVLSAALW
						SS*RGDSDSEGDLRATASITCRT
18492	48860	A	18599	1	2310	
18493	48861	Α	18600	3	1124	LYNRRRRRRCSHCRHRCRRLSS
						GLRKEEVISLGASLGRVFVPCSP
1						PTVSAARGPTGAPGGPNSKPLS
						GCCDDGFNLGRQQWGNPLPFC
		1				SKTISSSLHWTWSQVNLVEILP
						AIFSSFLNLQHVNLWFLAAAM
			į			KAVTEQGHELSNEERNLLSVA
						YKNVVGARRSSWRVISSIEQKT
						ERNEKKQQMGKEYREKIEAEL
						QDICNDVLVRGQCFCFEQWFL
						NSINLMYKCRLSNYFRYLSEVA
						SGDNKQSK*YL*KEIRPVMEPV
						FFHRDFLTILIRLGLALNFSVFY
						YEILNSPEKACSLAKTVRKTLC
						DI*P*QNSACVINFILFLNILRDN
						LTVSTTSTGFIVSFLFTYLIIHCY
						LQEGICTIKCSYSFKLLNLL
18494	48862	Α	18601	340	1323	AARPQKRHNNHSSLLTGSHIH
						WKRGRVQNIIADVLQFCLYLLT
						VLFVPCSPPTGMTMDKSELVQ
						KAKLAEQAERYDDMACSSERH
						VTEQG\HE\LSNEEKKSCSLVAY
						K\NV\VGARRFLPGRVISSI*A*K
						QRRNEKKAADGQKSTREKIEG
						RTCRTFCN*CFWELLGQISLFPN
						ATQPRKVKVFYFEK*KGDYF\R
	ļ					YLF*KWQS*RQQTQPTCVGTSP
				ŀ		ARAYPGSNLKI*LRKKMQP\TH
						PNSVLVLAPKFPQSFYYEI\LNSP
						\EKGLVSLGKNGHFD*SNLLNL
						DTLEWKSLIKDSTSDSWQLLR\
<u> </u>						DN\LTLWDIRKTRGDEGDAGEG
18495	48863	Α	18602	33	89	
18496	48864	В	18603	167	232	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	l e e e e e e e e e e e e e e e e e e e	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
18497	48865	A	18604	14	600	 MAELAPASPSDIKASVSNGDTT
						LLCSRRQSCGMNEVRQVSLTYP
						GSPAPSHSLPLQPRSGGSLCPSR
						AW/PDPHQLFDDTSSAQSRGYG
					1	AQRAPGGLSYPAASPTPHAAFL
						ADPVSNMAMAYGSSLAAQGK
						ELVDKNVSGRGWWEWGDARG
						HRASDLSSASPD\HRFIPITKLKY
,						YFAVDTMYVGRKLGLLFFPYL
18498	48866	В	18605	1	1230	
18499	48867	В	18606	41	931	DDDCDKCDI DDCUCCEVO ODDVI
18500	48868	A	18607	3	166	PRPFPKSRLPPGIIGGEYD\QRPN
					:	PILP\GRGGPNDRFPF\RPSRGRP TDGRLSFM
18501	48869	Α	18608	212	1881	CLRVLLLKRTWPVEVPETGARP
						/TGGILRSH\LRQSLLCT\WGYSS
						NTRFTITLNYKDPLTGDEETLAS
	i					YGIVSGDLICLILQDDIPAPNIPS
						STDSEHS\SLQNNEQPSLATSSN
						QTSMQDEQPSDSFQGQAAQSG
						VWNDDS\MLG\PSQNFEAESIQD
						NAHM\AEGTGFYPSRTPCSW*L
		1				NPVEGQVP\HSLETLYQLADCS\
		1				DANDALIVL\IH\LLMLESGYIPQ
		-				GT\EAKALSMPEKW\KLSGVYK
						LQYMHPLCEG\SSVT\LTCVPLG
						\NLIVVNATLKINNEVRKC*KGC
						SLLPESF\ICKEKLGENVANIYK
		1				DLQKLSRLFKDQLAHPLLAFTR
						QAL\NLPDVFGLVVLPLELKLRI
						FRLLDVRSVLSLS\AVCRDLFTA
						SNDPLLWRVFYICVNFRDNTV\
						RVQ\DTDWERTVQGRRAHTKE
		1	i			KESPKGRVLWMLLAIRQTH\TIP
·						FYSQPLWHPRGHFPKLPAFPPGI
			1			YSGGWNMDQR\PTFPYVGRTQ SSSLNSWVLGETPKPSFLPLRPR
			Ì			FDPVWPHFPGP*PPSLPRARAGP
						NDQISPFRPQQGSGQLIGRLSIH
18502	48870	Α	18609	1	1065	
18503	48871	Α	18610	1	1513	
18504	48872	Α	18611	1	2247	
18505	48873	A	18612	3	421	
18506	48874	Α	18613	1	103	
18507	48875	Α	18614	3	435	
18508	48876	Α	18615	268	501	QLECIPGTWELWKEEQERAMA
						GGGLKKIISATEKKLARMLLAP
						SASWCCSGP*VFHGST*WGKPG
				<u> </u>		PVSVSLLQGVLGK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18509	48877	Α	18616	3	296	EAMEAVILTCDPATPAASYQW
						WMNGQSLPMTHRLQLSKTNRT
						LFIFGVTKYIAGPYECEIRNPS*V
			}			PAASDPVTLESPPLLFNGLATTE
						TLRKRGNP
18510	48878	Α	18617	3	1464	GAVRTWGRGFQTEKCQASLLN
					<u> </u>	FWNPPTTAQVTIEAEPTKVSKG
						KDVLLLVHNLPQNLAGYIWYK
1						GQMKDLYHYITSYVVDGQIIIY
						GPAYSGRETVYSNASLLIQNVT
						REDAGSYTLHIVKRGDGTRGET
						GHFTFTLY\LIPWTLLYL*RSLA
1			į			QVSHYETPGKTAPPCGSTA**L
ł						N*PPDLTLFSPVLFLLKYPVPGQ
		1				AFQYPKGLKTIGSSITHL*DVLG
						KGSCRENIPRAAK*D*N*EDSST
ŀ			ļ			ACSK*GPQVETPKPSISSSNLYP
						REDMEAVSLTCDPETPDASYL
			ļ			WWMNGQSLPMTHSLQLSKNK
						RTLFLFGVTKYTAGPYECEIRN
						PVSASRSDPVTLNLLPKLPKPYI
						TINNLNPRENKDVLAFTCEPKS
						EN\YTYIWW\LNGQSLPGSVPR
						VKRPI\ENRVLIFTPCSRGIETG\P
						Y\QCEI\RDRYGGHPQLPSHPGM
						SSITTKHSGLYACSVRNSATGM
		<u> </u>	10/10			ESSKSMTVKVSAPSGTGHLPGL
18511 18512	48879 48880	A B	18618 18619	79 96	1516 1730	
18513	48881	A	18620	9854	11033	ESLCFPKTEQEASQSLSP*IIIHLS
16313	40001	A	18020	7654	11033	CDTCTSYWLSRTHVG*QVGDA
						NSE*RMPVEESKVPHRTIFCYPH
						S\KLPMPYITINNLNPREKKDVL
						AFTCEPKSRNYTYIWWLNGQS
						LPVSPRVKRPIENRILILPSVTRN
						ETGPYQCEIRDRYGGIRSNPVTL
		1				NVLCEYPLFLCGPGHQLNSKRP
		-				EARPLSLSPVQV*TTLLLDIRPG
						HDSLPWEILGRHSLNQEYKGRG
						ALVMGH*GPTACNERNRGIPQ
						AWA**T*MGFGCHLRLCLGSEG
						HCVPLRDQEHPLPLDDITCGFIL
						FPPDGPDLPRIYPSFTYYRSGEN
						LDLSCFADSNPPAEYSWTINGK
						FQLSGQKLFIPQITTNHSGLYAC
						SVRNSATGKEISKSMIVKVSGK
1	Ì					WIPASLAIGF
<u> </u>	<u> </u>	<u> </u>	L	L	l	WII ADDAIGI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
18514	48882	I IA	18621	3	1350	GRTAQLTAVLRKLLDPRLISTE
.001	1.0002		10021			ENTQAAETMGPLSAPPCTHLIT
		1				WKGVLLTASLLNFWNPPTTAQ
						VTIEAQPPKVSEGKDVLLLVHN
		1				LPQNLAGYIWYKGQMTYVYH
						YITSYVVDGQRIIYGPAYSGRER
						VYSNASLLIQNVTQEDAGSYTL
		ł				HIIKRRDGTGGVTGHFTFTLHL
		ĺ				ETPKPSISSSNLNPREAMEAVIL
		1				TCDPATPAASYQWWMNGQSLP
						MTHRLQLSKTNRTLFIFGVTKY
						IAGPYECEIRNPVSASRSDPVTL
						NLLPKLSKPYITINNLNPRENKD
						VLTFTCEPKSENYTYIWWLNG
						QSLPVSPR\VKRPIENRILILPNV
						TRNETGPYQCEIRDRYGGIRSD
		l				PVTLNVLYGPDLPSIYPSFTYYR
						SGENLYLSCFAESNPRAQYSWT
İ						INGKFQLSGQKLSIPQITTKHSG
i	-					LYACSVRNSATGKESSKSITVK
18515	48883	A	18622	273	2271	CGSWLMTQFSMDKRQGRSRSV
18313	40003	 ^	18022	273	2271	TITNTAVTSLEWVPFHAFAIPRS
	1					TLVSQQGATSHTDNTQEMDSP
						GQEPGFAKEVEVEPGFQPLPQM
ĺ		ļ				TLLVPAMCLLLHGACSAKGFC
ĺ						AAPHFLLASPMGKGQVPLNPFS
		l				FTLSEELDLPQSLKRNPKGCIAR
1						RAKPILAAERHKRLILHTSTKEN
			İ			TLLDNRVIECLTMEAVAKFNIM
			ļ			KEREGPSHQPALSPPKARWHIL
		l				WQKLMFTASLLTFWEP\PTTAR
		l				VTTEAMPFNATEEEEEEFFLLA
						HNLPQNLTRATIW\YK\GQMRH
1						LYHYITSYVVDGEIIIYG\PAYSG
		l				RETAYSNASLLIQNV\TRED\AG
						SYTLHIIK\RGDGTGGVTGRFTF
						TLYLETPKPSISSSNFNPREATE
i		1				AVILTCDPETPDASYLWW\MNG
						Q\SLPMTHSLQLSKANRTLYLF
						G\VTNYTAGPY\ECEIRNPVSAS
						RSDPVTLN\LLP\KLPKPYITINN
		ŀ				LKPR\EE*GCLRPFTCEPKE*GTT
						PYIWWVKRFRSLPVSPGVKRPH
						LKNRI\LIPTPVFTRKWKQGPYS
						MWNYGDRYGG\IRSLPSSPWNV
ĺ						LLWVQDLPRILPFHFTYYRSGR
		1				KSSYLSCSGGTLNPP\AQYSWD
						N*WKSFQLTRGQKLFYSGHITT
						N*WKSFQLTRGQKLFYSGHTT KHSGLYVC\SVRNSATGRESSK
						SMTVEVSDWDITLNSTSTSNSIF
18516	48884	A	18623	1	1131	OMIA DA OD A DLI TUO LO LOUOIL
10210	140004	I _A	10023	11	1131	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18517	48885	Α	18624	1	603	MDIADPFSGFPMGMGGFTNMN
10517	10005		1002	1		FGRSRPAQEPTRKKQDPPVTHD
						LRVSLEEIYSGCTKKMYQED\KI
						SHKRLNPDGKSIRNEDKILTIEV
						KKGISLPEALCG/TTVNVPTLDG
						RTIPVVFKDAIRPGMRRKVRGE
						GLPLPKTPEKRGDLIIELEVIFPE
				i		RIPQTSRTILEQILSSYLSYLSSP
						GLTRDLSRAQGFLDLSTSCGP
18518	48886	A	18625	3	1228	WSAAEPVDGDSGSAGRRRGPW
10310	140000		10023		1220	PKTTYHPWAVARGASEKEIKR
i						AYRRQALRSHPDKNKEPGAEE
						KFKEIAEAYEVLRDPRKREIFD
						HYREEALKGSGPSGG\SGGGAN
						ATSFSYTFHGEPLAMFAEFFGG
						RNPFDTFF\GQ\RNGEEGMDIDD
ŀ						PFSGFPMG/LWGGFTNVNF\GRS
ļ						RSAQEPARKK\QDSP\VTHDLRV
İ		1				SLEGDLTAAVPRRCTKKMKISH
						KRL/NTPDGKSIRNGDKILTIRSE
		1				RRGWKEGTQITFP\KEGDQTSN
1						NI\PADIVFVLKDKPHNIFKRDG
						S\DVIYPARI\SLREA/LCVGCTV
						NCPHSGTGRT\IPVVFKECYQGL
						GMRRKSSWENGLPLPQNTREN
						VGDLII*GFEVIFP\ERIPQ\TSRT
						VLEAGSFQYSYLEVPQGLTRGP
						FPELKDFWDLFLPVVGP
18519	48887	A	18626	321	1121	GDAVPSAPRRPSRRTPGCGSRG
10317	10007	'	10020	321	1.2.	ALQPGPPPRA\PPPEP/PPRGAAA
						AAPPAGPAAPAARQPPRRPPAQ
						PRSCPRRVRPTPAWSPPGRRGP
						RRSRRRPPGGPGPAAPLRLSRA
						HSPPAPPGSPYRPHGAGTSVAP
						WTRPPAARGTEADPGRCPVSAP
						GTAPGQMRGRGSATQRLHRGH
						STAPGAGGRCGRRPGGSGRPG
						AGAGPRRSTAAPPRCRRRSPTG
						PRSPRGPRSARPARGSVGGPRG
						TPGKQPGPRRA*TSGPEPRCCG
ļ						HARG
18520	48888	A	18628	191	653	KLEIQRLLRMTTECIRHTKQKK
10020	10000	ļ' `	1.0020			FGSPGLQPPLPGNAKSRADSTA
						LPALPAQQKETKQQEPGESS*E
						PGGSSPSRMERAEAAPAQS/PA
						GGTGTGQGWGAPHLGPALARR
						VTGPPQHGSPGPRTRPTPYPAR
						C/PSRPIFVPAIVWSRLRAPPGLP
18521	48889	A	18629	141	319	ON SKI II VIAIV WSKEKAI POEP
10321	70007	147	10029	11-11	313	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
18522	48890	A	18630	178	722	PLGNVSCSHRSFQTARVDYSLI YVLAWGQKKEVWELRNWQTH ALMKLKNKLREPRTSVNKD/SP KSLLYSCSYSYFDEPVELRSSSF SSWDDSSDSEYETHLLHLKLV* PNLAVFNCRPTARRKPDYEPVE NTDEAQNSFCKTAHNLWSLTFP FPCLL*YETRARLESWSLSCLR YEHISSG
18523	48891	A	18631	2	367	
18524	48892	A	18632	1	381	QRYSSRRATPRHIIVRFTKVERK ETMLRAAREKGRVTHKGKPIR LTVDLLAETLQARREWGPIFNI LNEKNFQPRISYPAKLSFISEGEI \NPQMLR/DFVTTRPTLRDLLKE ALNMERNNRYLPGRPL
18525	48893	В	18633	1	846	
18526	48894	A	18634	3	1170	VGSMTGEASGNLQSQRKAKRK EARLTCLEPVEEPGSGTHPPMC YLAPPMWKCQKPETGIKEGDR DHNSLPAREENQTENESDELTE AGFRRWVITNSSRLKEHVVTQC KEAKNFEKSLKELLTRITSLEDN INYLMEPRNTAREHREAYSSIN SQIDQAEEKIPEIEDQHNEIKRE DKIREKRMKRNKQSLQEIWDY VKRPNLSLAGVPESDEENGIKL KNTLQDIIQENFPNLAKQANIQI QEI\QRTPQRYSLRATPRHIIVR FTKVEMKEK\MLRAAREKGR\V THKGKPIRLT\ADLSAETLQAR/ RTEWGPIFNILKEKNF\QPRISYP AK\LSFISEGEIKYFTDKQMLRD F\VTTR\PALKRAPGKEGT*TLE RDN\RYQPLQKHAKL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18527	48895	A	18635	173	1504	QDLRLFYLKDTMFSKLAHLQR
10327	10075	``	10033	'		FAVLSRGVHSSVASATSVATKK
		1				TVQGPPTSDDIFEREYKYGAHN
		1				YHPLPVAL\ERGKGILLMGM*E
		1				GRKYF\DFL\SSYSALSNQG\HC
						HPKIVKCSGRVKWDKIDL*HLE
						AFYNNGTLVNMREYYY*TFST
						YHKVLP\MNTGVEAGETA\CKL
		1				A\RKWGL/YTVKGIQKYK\AKIV
						FAAGNFWGRTLSAISSSTDPTS
		1	ł			YDGFGPFYARDFDIIPY**SCPH
			ł			MEQCSFKDPKCGLAFHG*EPIQ
						G\EAGV\VVPDPG\YLMGSAESS
						GTRDQVLFICLMEIQTGLARTW
						*DGLAVDYENVRPDIVLLGKAL
						SGGLYPVSAVLCDDDIMLTIKP
						GEHGSTYGGNPLGCRVAIAAL
						EV\LEEENLAENADKLGIILRNE
						LMKLTLP/SMLVTARKKEKGIY
				ł		LNAHLAIKGNQKIWDA/WGKV
						CLRLQDNGLLPKPTSVATLSRF
ļ						APP\LVIKEDELRESIEIINKTILS
18528	48896	A	18636	2	348	
18529	48897	A	18637	588	801	RTAYFCQYHTASVYSERAMPP
				1		GCPEPSQA*ELQETGHRQVALR
				1		RSGRPPKCAERPGAADTGAHCT
						STDGRL
18530	48898	A	18638	175	510	NSESPGPLWWLSCDNPAKDRIY
						STEADGLGLPDQSFVVHRAQL*
						CVAG*LGTHSRFQEWAV*D*SD
				1		LLKAQLHFKVGRCVWEAGGGL
						EWVLGGRCRCRVPQPLETLIQ
				1		WGPW
18531	48899	A	18640	1	1431	
18532	48900	Α	18641	202	994	RWQGQRLLLDSNIRLWVVLPIR
						YSFTFFRRALIRHYV\SILLAERQ
						RSFTQGNKYLDSSSPNFEGRV/R
						FRGKLGNYIPKQSFFG/RRKFLF
						FPNPEDGIFSKKLKREVVPPSPM
				1		TDPTMLTDMMKGNVT\NVLPM
					1	ILIGGWINMTFSGFVTTKVP\FPL
						\TLRF*A*WLQQGNRSYSPFRCI
						PG*/WFCIPGYFPQCILGFRSI/YT
			1			LWISGGKDNGRWNQSRMM/DR
				1		SRMTGASHGPWPADSGRRFPN
				1		KRWEAL\ELTDHQWALDAAGE
						DAA
18533	48901	Α	18642	97	224	LPTKSTWVAGKK*KKRSNIKD
						YSTPLTSYRSIRSHAFKIVAS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	la contraction of the contractio	*=Stop codon, /=possible nucleotide
ĺ	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		İ		sequence		
18534	48902	Α	18643	117	457	IWGDVEKGK\KVFYL*RCSQCH
10331	10502	1	10015			TV*/ERGGKHKTGPKSP*VSFGR
						RTGQAPWDTLYTAANK\NKGII
1						WG\EDTLM\EYLENPKKVHPLD
1						TKMIFVRHLRKREERADLIALS
						SKKLLN
18535	48903	A	18644	411	1054	VTGAQRLVTCRMGKQNSK\LA
10333	.0500	-				P\EVMED\LVKSTEFNEHELKQ
						WYKGFLKDCP\SGRLNLEEFQQ
						LYVK\FFPYGRRLPSFAQHAFPN
						LSTRMGDGTI\DFREFICAL\SITS
						RG\SFEQKL\NW\AFNMYGPGW
						VMGKITRVEML\EIHRGLSYQN
					1	G*GTVIMMKMNEDGLTP\EQRV
						GQDFSARWVRNKDDQIYTGID
						FKGSCQRADPFIV*LL\QCDIHK
18536	48904	Α	18645	3	475	GAEGARGGGSSYSEMAETVAD
				ļ		TRRLITKPQNLNDAYGPPSNFL
						EIDTNLPIFKLKESTVRRRYSDF
						EWLRSELERESKVVVPPLPGKA
		İ				FLRQLPFRGDDGIFDDNFIEGKK
						TRGWEQFINK/VSLGHPL\AQNE
						PLFFHMFLQDEIIDKSYTPSKIR
18537	48905	Α	18646	1	695	LRARSLRDRCARAPCPHGGQQ
						RRRRLNAEGAEGARGGGSSY
						SEMAETVADTRRLITKPQNLND
Į.						AYGPPSNF\LEIDVSNPQTVGVG
	1					RGRFTTYEIRVKTNLPCFQS*KN
						LIVRR\RYSDFEWLAKWNLERE
						SK\VVVPPLPG\KAFLRQLPF\RG
						DDGNILMDNFI*/EERKIRGWSS
	1		1			FLNKVA\GHPL\AQN\ERC\LHM
						VFYQDEIIDKKLLLPSKNKAMP
		L				WKFGPRKGAKNVDY
18538	48906	С	18647	3	474	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18539	48907	Α	18648	156	1254	RPLHTLEVLRRMTKRLVTKAA
						ASAARAQPAEPRDRKYDKHLS
						YLATWGRSAGRAGSPRGARGR
						ELARAPVSLPAGVPGRGSAAG
						HSSEARTGPNGKMGGSSREPGT
						EKVPCPRGLSPTREVGHQVHGT
		ŀ				LGAGPNELHPSGPAGARCRPGR
		İ				REAKLGDAARSEAQSSAHKFHI
						DSLSLPWNPTAMEMLMPKKNR
						\IAIYE\LLFK\EGVMVPKK\DVPI
						PKHPELAEQKCAPTFHVMK\A
						MQ\SLKSPRPT*KEQFAW\RHFY
			İ			WYLT\NEGIQ\YLRDYL\HLPPEI
						VPAHPYGRSPSRRLARPRPKKV
						WKGEATLRDFTRGEADSFLCH
						S/DSYRRSAVPP\GA\DKKAEAG
		1				GLGQKPNFQF\RGGFVRGRGQ\P
18540	48908	В	18649	1	1053	
18541	48909	Α	18650	1	529	MGRYMEREPLQGNHRRSGLSG
						ERCSHMGGVLIERRELQTETRK
						EGVVEWCWLEKEVLATITKPV
	1					RGDKNGGTQVVKHRKMPRYY
1					,	PTEDVPQKLLSHG\KKPFSQHM
						RKL/FLYRRHARNLSLPPQPELII
						SDVK/IPRHPADA*LEEQQLSEP
						E/HSEGTSY*RIPEKVLSYPAIRP
18542	48910	В	18651	258	363	
18543	48911	Α	18652	197	398	ARMSQEKDFHKVMSALKARTG
		1				HLHFFCGGMSSVKVGQGFSLLL
						*FFSYFRPSGCILPL*ELPEARCP
18544	48912	Α	18653	215	932	GTQSCNQKVLNSANNLQELGIG
						PKFQMRLKPQLAPMQPNRGTS
						LGRSIPCPPVLCSMRKIHLRPQV
						LRPTSPRNISPILNRVSEVSDHA
						GTPALVLHP*RQVPLFWGRGK
			ľ			YPNPFSLCLYPFSAFLGGKKHP
						TPSPSP*VASPLF*RSKYPNLVSL
						CPSPLFPRPDLLSLWPNPLF\RTP
		1				TSYISVPQSLISMPRPLISAPQPLI
		İ			ľ	STTQPLSRFSGRSLLQVPEIWPP
						GQGMPAARDSS
18545	48913	A	18654	1	378	
18546	48914	Α	18655	14	380	
18547	48915	C	18656	228	260	
18548	48916	С	18657	99	335	
18549	48917	Α	18658	55	474	RQVPLFWGRGKYPNPFSLCLYP
						FSAFLGGKKHPTPSPSP*VASPL
		1				F*RSKYPNLVSLCPSPLFPRPDL
						LSLWPNPLF\RTPTSYISVPQSLI
						SMPRPLISAPQPLISTTQPLSRFS
			}			GRSLLQVPEIWPPGQGMPAAR

SEQ ID	SEQ ID NO:		SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
18550	48918	Α	18659	1	187	AELATGNVKVEASGTPEEKAG
						SVLG*ACFAGP*ERCCWQNN*D
		<u> </u>				GIISIQSYCLCNLGCFIQT
18551	48919	С	18660	1	357	
18552	48920	Α	18661	3	148	
18553	48921	Α	18662	1	312	
18554	48922	A	18663	149	457	
18555	48923	В	18664	1	675	<u> </u>
18556	48924	С	18665	110	928	
18557	48925	A	18666	495	817	LTSPSRCTIIEKSCSSLPPL*DKP
		ŀ				QPHLQHTRTSKRLNCSSQVFLQ
						NLLPEELATSSRNLATGPRNAC
		ľ				SPGFLLSHIPSVWDPTGNRTVQ
		ļ				LTWQPLPEPLELWPKAL
18558	48926	A	18667	228	3021	VFDRTQNVQGRIMEAPEYLDL
]				DEIDFSDDISYSVTSLKTIPELCR
		İ				RCDTQNE\DRSVSSSSWNCGIST
						LITNTQKPTGIADVYSKFRPEKR
		ł				VLPLKHQPETLENNESDDQKN
		1				QIVGEYQKGGESDLGPQPQELG
		ł				PGDGVGGPPGKSSEPSTSLGEL
						EHYDLDMDEILDVPYIKSSQQL
		ŀ				ASFTKVTSEKRILGLCTTINGLS
		ŀ				GKACSTGSSESSSSNMAPFCVL
		1				SPVKSPHLRKASAVIHDQHKLS
		Ì				TEETEISPPL
18559	48927	Α	18668	1440	1764	ERSTYNLRSSDQP/RPRNILTNF
						KS/VDKGDTFYPWTHNSGASH
						GLGR\RLPWC*ELATSARNLTP
		ł				RPRTAGSPGFLLSHVPSVWDPT
		l				ANRTVQLTWQPLPEPLELWPK
18560	48928	Α	18669	1	3255	MLTLTGAGGESWIIRPGNLLEQ
ŀ						LGAEETLGRRQEPGDEVTAHFL
		1				LQGVINMGPVPYPGAHSSNMP
İ						GSVSPILGHSVKLFEEYYPVGD
İ						PLMGSGRPHTPRKKIWSDLNA
•						HSDLYLALDWTPLPEQSEPGPG
						LAAWKPFQAEEYTCKCAEGNE
						NTPFDLRKGTLPCPRNLERARQ
						AYMHIEEKFPFELTEQHICEFQS
						MCYVLVRVLGACSQSTWLILEP
						DGVSGPSKGASEAIGQCQSSAA
						KPRRSGKESVREPWA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18561	48929	Α	18670	1	1442	CSEYEDSSPAPVPATDLSSTLSS
			•			SVPQPQDTGTSQQLHPLDPWHE
		l				LLRAQELQGATNHKGYSHAEH
		Ì				EHAGLGVQGGNGALAFSNSGH
						RHAVPTISSGTGRRRTPSSSAFG
						LLNLHQWFVSGFQAFSDRLKA
		ļ				ALSASLLLRFGDSDWLPSSSAC
		i				KCLMLGLHFVIVGNICATLKEK
						YSSMLHLDVTMKKNGEKRTRL
						QKRKKGMPPHPA/S*GPKHSCH
		1				HSPCQCCPSSAK/SASELQGSWT
						PFGGLLTRTRMKFGAKTRDLD
						QGTSLGRSLPCPPTLCSMRKIHL
		l			İ	RPQVLRPTSPRNISPISNPRQRR
						QVLSMDPKLRHRSRTGK/DSLP
						LVFNHCRDTSLIIHPCFKGVRPR
						RDACLGPSPLAASPAFLEKGQD
	<u> </u>					LINLAFKVYNNRKKLQFLASTV
		1				RQTAATSPAHKNFQMPEPQRP
					1	GVPPEPPTGACYMCRKIWPLG
						QANARSPGFLLSRVPSVRDPTG
10550	10000	ļ	10671	1.40	1005	NRTVQLTWQPLPEALELWPKA
18562	48930	Α	18671	140	327	WELL COLORS OF BORDS
18563	48931	Α	18672	95	115	IIRHLCND*TPREGCLPSP*PAW
		<u> </u>			 	SDTFETWVNNQASLQ
18564	48932	Α	18673	218	674	MPISRPTARFKRIKVYYHSPAT
						AWPSKAYKLPLQFPHFTCPKTR
						QGLQVTSGSAPYQPNCFVYPPR
						VAKTKYSPILNTSLHNPLLCSGS
		3				QTCFLY/SFLCTFHPSLSSLSLGL
						TLTPIRLSKLRGLYCHKASQTA
						PITSIKPKFLPHLLPISA
18565	48933	Λ	18674	1	268	
18566	48934	Α	18675	1	458	
18567	48935	A	18676	3	461	LLLTQSLFGGLFTRTRMKFGAV
						TRIGGPPLGNQSPSSCSLLHEKD
						PPTTSGPQTDQPKKHLTNFKSA
						ARPTFLGQGQVPLNPFSFTLS/E
						QVLLS*AARTPQSLISTPQPLISV
						PQSLISVPQPLLYFSGGQEPPPPP
						LLCVSSLFSRLASFTM
18568	48936	С	18677	156	329	
18569	48937	Α	18678	449	667	
18570	48938	A	18679	79	311	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18571	48939	Α	18680	1	693	MAPTNCKGSRKCGRPHEQMPH
						PYLPLLTLFSDSARLHPGEINSR
						VPHTKPVWWSLHTDAYEIWYH
						DSDWGDLPWEINPLSSCSLLHE
						KDPPTTSGPQTNQPKEHLMNFK
						SGPHWKSDCPTRQPLPGPLELW
		1	•		:	PKA/HLTDSFPDLLSLAAED*CC
						LIALEAPWTITDAELWITLTVED
					•	SQLYEDTLAGRSVLIKNLTP*TL
						Q/PLDGLDPT*SSIVPQLPSACRI
						LPTGFTVPE
18572	48940	Α	18681	328	865	
18573	48941	A	18682	451	1238	TALLLTQSLFGGLFTQTRMKFG
	·					AVTRIGGPPLGDQSPVLLFFVPR
						ERSTYDLGPQTDQPKKHLTNFK
						STSFVFSSCIPPP*PSSISLLPPW/T
					i	TDHAPLTISL/TT*SPLPCSMPIS/
					<u> </u>	ASHSML*KD*SLLSLACYSMAF
					[*SL*TLLTIPPFYLS*NQRRFTG*
						FRICALSTKLFCLSTPWCQTHIL
						SYPQYLPPQSIILFWISNVLSLLF
			į			LCTLHPSLSSLSLGLTLTPFRLS
						KLPGLYCRKASQTAPITSVKPK
						FLPYLLPISA
18574	48942	C	18683	870	1037	
18575	48943	A	18684	933	4323	AASHLCGTPLEIGLSNSPGSHSQ
						SPWNSGPRLSDCFPDLLG*VAE
						D*HCPIASEASWTVT\ELWVTLT
						VEVAATALILLEALKITSYAPLT
				ļ		LYSSHNFQNLFSSSHLT/PYTFC
	ļ					PQ/GPFSYTHSLLSLPQSPLLLA
						QTSIRPPTLFLIPHLTPMTVSL*S/
						ILTFIPFPHISFFPVPHPDHTWFID
						GSSTRPNRHTPAKAGYAIVSST
						FIIEATALPPSTTSQQAKLIALTQ
						ALTLAKGLLVNIYTDSKYAFHI
18576	48944	A	18685	849	1944	WQVPLFSGRGKYPNPFSPCLYP
10370	10777	'`	10003	1377	1777	FSAFPGKRQELATSARNLSDHQ
		1				AKECLQPRIPPKPCPIFAGPHWK
		1				SDCSTSPGQPLPEPLELWPKA/H
						LTDSFPDLLGLAA\ED*HCPIAS
i		1		1		EAPETITDAELPVTLTVEGKSIP
						CLIDTGATHSTLPSFQGPVSLAP
						ITVVGIDGQASKPLKTPPLWCQ
						LGQHSFMHSFLVIPTCPLPLLGR
		1				NILTKLSASLTIPGVQLHLIAAL
						LPNPKPPLCPLTSPQYHPLPQDL
	<u> </u>		L	<u> L</u>	<u> </u>	LENEAL FLORE 1 SPQ 1 HPLPQDL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
		l		sequence		
18577	48945	Α	18686	2	377	KGATIVYALDTLFIMEMKHEFE
		1				EAKSWVEENLDFNVNAEISVFE
				}		VNIRFVGGLLSAYYLSGEEIFRK
						KAVELGVKLLPAFHTPSGIPWA
l						LLNMKSGIGR\NWAWASGGSSI
				1		LAEF\GTLHLEFMHL
18578	48946	Α	18687	215	1135	PITLRRHLFFSGIGRNWPWASG
		ļ				GSSILAEFGTLHGMKIVF*/HFH
						GNILFSI\KVMNIRTVLNKLEKH
						YLN*KLNHSKIHFPLFPDHVSV
						GGLGDSFYEYLLKAWLSINLRR
						DYKMLKNKCFSQAIETHLIRKS
		ŀ				SSGLTYIAEWKGGLLEHKMGH
						LTCFAGGMFALGADAAPEGMA
		l			}	QHYPVSTVQENNPYSFIPCPLV
l						MKLGPEAFRFDGGVEAIATRQ
l						NEKYYILRPEVMETYMYMFRL
ŀ						SGSDFFSSFLLLPPQASSCFQTLF
						DLFKVPDSLCSILEFRRHSSRQP
						YSTDQPFIYLVGKRILFCPLT
18579	48947	С	18688	304	400	
18580	48948	Α	18689	16	450	
18581	48949	Α	18690	1	777	GHSMDMRVPAQLLGLLLLWLP
						GARCVIWMTQSPGTLSLSPGER
						ATLSCRASQRVNSNYLAWYQQ
						KPGKTPKLLIYGASNLETGVPS
		}				RFSGSGSGTDFIFTISSLQSEDIA
						MYYCQQYNNWPHPS\TFGQGT
						KLE\IKRT\VAAPSVVHLPGPSD
						EQLKIWELPSACVPA*NNF\YPQ
						SRPKVQWEGGITPLQ\SG*LPQE
		İ				SVHRGRTRQGTAPYSPQQAPW
						TVRQSRITEETQSLTACEV\TPS
			_			GALRFRPVHKGAFNRGESF
18582	48950	Α	18691	3	688	HASADAWAANAEEHRHDRPR
						GTLR\EYKV\VGRCLPTPK\CHT
						PPLLPACEIFAPNHVVAKS/RAF
						WYFVSQLK\KMKKVFQGKIGL
						LVGQVF*ESSPLR/V*KNFRGSW
						LAAMDSRERAPTNMLPGNNRG
						P*PPAGGCSTQLLTRDNGVAPG
						TGAPKPHFHFRFIERLEGRFAG
						QQSCRRP/ALVQSSFH\DSKIKFP
						ACPNGVL\RRQHKPRF\TTKRPN
		<u> </u>				TFFLGAGPSSGVCPK
18583	48951	Α	18692	1	784	

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO:	Nucleotide location of first		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
18584	48952	Α	18693	1	4713	MQGPYLLLWLPDSLEEGKEGL
	į			-		YLGNCIHTCLVTVPEDAFFPHF
						LTQHSGTHLPPSSDIPSHFMNNS
						TGRLAQRILGTSTEAFLSPEVM
						VFVHARNATVRTAMSLIERAK
					•	NCGHIPFFFPTQGHDYVLAEKQ
						VQRSRNKQVRELFPDGFSIHHA
						GMLRQDRNLVENLFSNGHIKV
						LVCTATLAWGVNLPAHAVIIKG
						TQIYAAKRGSFVDLGILDVMQI
						FGRAGRPQFDKFGEGIIITTHDK
						LSHYLTLLTQRNPI
18585	48953	Α	18694	313	489	SSCFKLHFQQLFKNCNTQMRN
						QIGFLRF*NNLVMLLSSSSVTNQ
	ļ					LELQKKIILLTCKMQ
18586	48954	Α	18695	1	851	ACGWALAQRPARRAMVAGID
						AGRS\LGVLSVVCLLHCFGFISC
						FSQQIYGVVYGNVTFHVPSNVP
						LKEVLWKKQKDKVAELENSEF\
						RAFSSFKNRVYLDTVSR*PSLST
						T*TSSR*KMKYEMESAKIVTD\T
						MKFLPYVP*VSFQSPTLT\CALT
						NGSIEVQCMIPEHYNSHRGLIM
	1					YSW\DCP\MEQCKRNSTSIYFK
						MENDLPQK\IQ\CSLSNP\LFNTT
						SSIILTTCIPSSGHSRHRYALIPIP
						LAVIPTCIVLYMEWVF*KCARN PDRTQLQLIGNRRMKTTA
18587	48955	C	18696	35	181	FUNTQUQUIONNOMETTA
18588	48956	A	18697	75	344	PQVLRPTSPRNISPILNRVSDHA
10300	70750	``	10057	/3		GTPALVLHP*RQVPLFWGRGK
			ł			YPNPLYLCAPIPYFRT/HNLISLR
						PNPLCPHPDLVSLCPDPFPAFLE
18589	48957	A	18698	1	268	
18590	48958	Α	18699	39	376	QYISELQFLASTVRQTPATSPAH
						KNFQTPEPQQPGIPPEPPPPGAC
						YKCWKSGHQAKECLQPGIPRK
						PRPISGSHSQSPLELWPKA/HLT
ŀ		ł				DSFPDLLGLAAED*HCPIASEAL
18591	48959	Α	18700	2	295	CQTTQGRLLTAGTPL*SFTHVS
						RVSDHAGTPALVLHP*RQVPLF
		Ì				WGRGNHISGTQELPNT*TAV/V
		1	1			QAF/LPEPPPTG/CLLHVPEIWPL
						GQGMPAGQDSS
18592	48960	Α	18701	81	402	VKFGPEIWCRDSDQGRGGVGT
						SLGRSIPCPPALCSVRKIYLRPL
						VLRPTSPRNISPILNRDPTVQLT
						WQPLPEPLELWPKA/HLTDSFP
						DLLGLAAEDRCCPIASEAP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18593	48961	ĪA	18702	165	439	PPRQAKMQNLAAPGSHSQSPW/
10333	10501		10702	103		TLRPKAL*LTPSQIFSA*RLKTD
						TARSPRKPPSFQGPVSLASITVV
						GIDGQASKPLKTPQLWCQLRQ
						YSFK
18594	48962	Α	18703	284	511	IILVVKWANGLRCLTSRHSFTH
					,	QSLPS/PLCPVQLVPN\SRLSPQS
					}	QPQASLSLSNLPFLQTHLTSPFL
						PRLLLARPK
18595	48963	C	18704	414	557	
18596	48964	Α	18705	1	1284	MNFEGTETLHCNGHCKVIAMT
					}	LGNQEVNQRNFSTGTTGPMEG
						MWLLSSNSHHSLKYAGALAILS
ļ						AFQYQERGSQAGHRDSWPETK
						ETHFICGPKTPALVTDWEGSLP
						LVFNHCRDASLIIHPRFSGVRPR
						RDACLGPSPLAATPQTITDTEL WVTLTVEVWDIST/LFPGN*SH
						A/TLPSH*NLITLTPLNANI/AIPQ
						HALKGLKPVITRLLQHGLLKPI
						NSPYNSPILPVQKLDKSYRLVQ
						NLRLINKIVSPIHPVVPNPPAFTS
					:	QITQAVSQALGIQWNLHIPYHP
						QSSGKVERTNGLLKVHLTKLSL
						QLKKDWTVLLPLALLRIRACPR
						DATGYSRFELLYGRTFLLGPNLI
						PDTSPLGDYLPVLQQARQAAN
						LLLPTPDPQPHEDTLAGRSVLV
						KNLTPQTLQPRWTGPHFIIYSTP
						TAVCLQDPPH
18597	48965	В	18706	1	933	
18598	48966	Α	18707	3	435	TKETGFIHGPKTPAPVTDWEGS
						LPLVFNHCRDTSLIIHPCFKGVR
						PRRDACLGPSPLAASPAFLEKG
						QDLINLAFKVYNNRKKLQFLAS
						TVRQTAATSPAHKNFQMPEPQ
						RPGVPPEPPPTG/CLLHVPEIWP
10500	19067		19709	2	111	LGQGMPAARDSS
18599 18600	48967 48968	A B	18708 18709	2	595	
18601	48969	С	18710	1	1344	
18602	48970	A	18711	227	385	VPILPQPLLLHPAIFLSPPLLTPG
1.0002	1.05.10					PAYSFVP*LALPHLPSNLLLKR WLEPKA
18603	48971	С	18712	148	258	WEELER
18604	48972	A	18713	215	693	
18605	48973	C	18714	1	816	
18606	48974	В	18715	1	513	
18607	48975	В	18716	1	1452	
18608	48976	A	18717	1137	1303	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18609	48977	A	18718	2	12107	IDMIFTPGPPSTPKHKKSQKGSA
18009	40311	Δ .	10710	[2107	FTFPSQQSPRNEPYVARPSTSEI
					,	EDQSMMGKFVKVERQVQDMG
						KKLDFLVDMHMQHMERLQVQ
ŀ		İ				VTEYYPTKGTSSPAEAEKKEDN
	l.					RYSDLKTIICNYSETGPPEPPYSF
}						HQVTIDKVSPYGFFAHDPVNLP
		Ì				RGGPSSGKVQATPPSSATTYVE
		ļ				RPTVLPILTLLDSRVSCHSQADL
į						QGPYSDRISPRQRRSITRDSDTP
						LSLMSVNHEELERSPSGFSISQD
						RDDYVFGPNGGSSWMREKRYL
						AEGETDTDTDPFTPSGSMPLSST
l						GDGISDSTVPLFLSSEILQKQVG
1		ļ				QSITSMLGFLSRGPSMKLCMGL
	1	l				ACVLSLWNTVSGIKGEAKKEK
						GMTFLPTTDSKKFFSLLSVTSYS
						SFAFHKFSVAVYNISNLKTVDP
İ						AKFPTRYCYCLNNRTNDLSDFT
						ALLVDIIGNSTSYLTEIFKSTSIL
1						SVNQSNESDCIFICVMTGKSGR
]				NLSDFWEIEEKYPIINYTFTSGL
						SGVLALLLTQSLFGGLFTRTRM
		l				KFGAVTRIGGPPLGNQSPSSCSL
						LHEKDPPTTSGPQTDQPKKHLT
						NFKSAARPTFLGQGQVPLNPFS
						FTLS/EQVLLS*AARTPQSLISTP
1						QPLISVPQSLISVPQPLLYFSGG
,						QEPPPPLLCVSSLFSRLASFTM
1				İ		GAFTHGTQTPSPTKATAPRYPQ
						TGDLSAEWPFTAGEEPVLVPRP
18610	48978	Α	18719	828	1063	RHSQAAED*HCPIALE/VPQTIID
						AELRVTLTVEGYSLPPGCLRGL
						MLSVSGFSRSTVQAVSGSTVLG
						SGGQWFSCASV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				Sequence		
18611	48979	A	18720	183	1140	LGSGDLPWEINPLSSCSLLCEKH
						PPTTSGPQTDQPKKHLTNFKSG
						ETKEMHFIRGPKTPVPVTDWEG
						SLLLVFNHCRDGLSDHSATFQG
						CQT/MAGMPALVLHP*RQVPLF
						WGRSKLLLARLS*VPILPQPLLL
						HPIILLSPPLLTPGPAYSFVP*LS
						MSIHQVVLRPDEQGPGQEQQPP
						LYVMEPGEINSFIAHTKPVWWS
						LHTDAHEIWCHDSDRGTSLGRS
						IPCPPALCSVRNIRLRPQVLRPTS
		-		İ		PRNISPILNQERQRRCILFVDPK
1						LRCQSRTGKAAFSWCLIIAGTA
						SLIIQPHFKGVRPCRDACLGPSP
						LAASPAFLGKEQAAPCQAELGP
						NSSSASAPPPYNPFITSPPHTRSG
						LQFRSVTSPPPPAQQFTLKKVA
		j				GAKDIVKASLAPALAQGVPGTP
	1					QATASEGASLKSWECLCGVNL
						ATAQSARVKEACQEACQSLPRF
						QRMYEKAWVPKQKTTAEVELS
				į		QRASTRVVLRGNVGLESPSRVF
						SIALSSGAVGGVHCPSDHRMIE
						PAAFNLSMKNPQRQSCLRA
18612	48980	Α	18721	372	906	LRSADLPWEINPLSSCSLLHEKD
						PPTSSGPQTDQPKEHLTNFKSE
						KKETRFIRGPKTPAPVMD*GRQ
						PSLGV*PLQGCLSDYSPRFQRC
						QTTQGHLPWSFTLSSKSHFSGG
						RGKSLLQVPEIWPPGQGMPAA
						QDSS*AV\PICAGPRWKSDCPTH
		-				LAATPKAPGTLAQGSLTPSQIFL
18613	48981	Α	18722	983	1980	KFGLVQLTLGKPLPEPLELRPK
						A/HLTDSFPDLLGLAAED*HCPI
						ASEAA*TITDTELRVTLTVEVW
				1		DIST/LFPGN*SHA/TLPSH*NLIT
				1		LTPLNANI/AIPQHALKGLKPVIT
						RLLQHGLLKPINSPYNSPILPVQ
						KLDKSYRLVQNLRLINKIVSPIH
						PVVPNPPAFTSQITQAVSQALGI
	1					QWNLHIPYHPQSSGKVERTNGL
				1		LKVHLTKLSLQLKKDWTVLLP
	1					LALLRIRACPRDATGYSRFELL
						YGRTFLLGPNLIPDTSPLGDYLP
						VLQQARQAANLLLPTPDPQPHE
						DTLAGRSVLVKNLTPQTLQPR
						WTGPHFIIYSTPTAVCLQDPPH
18614	48982	В	18723	1	271	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18615	48983	A	18724	803	1158	CSWHDRFPDWKAGNLLSLAA*
	·	1		į		D*HCPIASEAP*TITDAELQVTL
						TVEGSGDLPWEINPPSSYTLLCE
						KDPPTTSGPQTNQPKKHSHQFQ
						IRDKGDTFYPWTHNSGASHGL
						GRQPSLGV
18616	48984	Α	18725	1855	2506	GASPSVWDPTENRTVQLHLGS
		ŀ				HSQSPWNTGQRSRDLTSFPDLL
						GLAA\ED*HCPITSEAPLTITDAR
						AMG*LSTVEGKPVPFLINTEAT
	1					HATLPSFQGPVSLASIT\VVGID
				1		G\QAF*TSLKLPNSWCQH*TIRR
				1		FKHSFLVIP\TCQVPLLG\EDTLT
					}	KLSASLTIPGLQLYLIATLLPNP
						KPPLCPPLVYPHLNPQV*DISTP
						SLGDRSCTPYHLIKT
18617	48985	A	18726	2	979	TSTMAVGKNKRLTKG\GKKGA
		İ				KKKVVDPFPR\KDWY*RRKHPL
						MFH\IRNIEKDVGSPRTPRDPKL
						QSDGLKGSVCLKVSPAELPE*L
						KFAFRKFK\LITEDVSGVKTCPD
						LTSHGAWDLYP*QNCCSMVQK
		1				MGRPMIEA\HVECSRLTD\GYLL
						\RLFCVGFYLKNRNISDHGRPSY
		-				AQPPNRVR\QIR\KKMMEIHDPQ
						RLQTNVP*KEVVNKLIPASIGK\
						DI\EKACQ\SIYPL\QNVF\VKKV
						KMLK\KPKF\ELGKLMELHGEG
						SSSGKATGDETGAKVERADGY
						GPPVQESVGKFRLPIVGNRSGH
		i				CVKKEKQSSRRRDFGVSVHPRI
18618	48986	A	18727	li	615	ASTAAPRMLLLFQPRYRTLQPQ
						RLLMPKKNR\IAIYE\LLFK\EGV
						MVPKK\DVPIPKHPELA\DRNVP
						NLHVMKAMQVSQSPRG\YVK\E
						OFA/WKKISYWCLTN*GYQGVS
						ISSRDYL\HLPP\EIVPA/TPLRRS
						RPED/SGRPSA*KGLEG*SDLAR
1		1				LHKNGEAGQRLPYRR\SAVPP\G
						\ADKKAEAGGLGQKPEF\QF\RG
						GFVRGRGQ\PPQ
18619	48987	T _A	18728	3	395	SAEVGAAETTLTELRRTVQSLEI
10019	10,0,	'`	10720	ľ		DLDSMRNLKASLENSL\GILLHL
						ESELA QTRAEGQRQAQEYEAL
						LNIKVKLEAEIATYRRLLEDGE
				:		DFNLGDALDSSNSMQTIQKTTT
						RRIVDGKVVSETNDTKVLRH
			<u> </u>	1	L	MA VOOR V VOETNOTR VERT

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
18620	48988	A	18729	323	1973	PCMWIGFAAVRKGLKDSSSASC GCTRRVRGTWWQGLSRVCVA RNEEVRLWGLCWLGSSPPAGW NNVFYMLISADVLACLEPRAN KEAANELAGAVDLGWMANVR RNPRDMVTSTHLGTHFHLRKTP PRGEKTCELASQQPLIEATTKT KSVAVSHILTYTPPTLNGVAAV SSRVHFSFWPPTLQTGLEQGSM SFTICFTTSTNYQSLHSIQPHSHS VQFISSTAKVYAVLRGLGSRTS VSFSTSFWSGWGSGGLAAGMP GIDNARLAADDFRVKYKTELV MCLSVESNVRGLHKVTDDTNV TRLQLETEMEALKEELLFMKK NQEKEVKGLQALIASSELTMEV NAPKSQDLSNIMADLPAQYDEL A\RKN*EELDKYWSQED/IEEHT TVVTNTSLLEVWSC*DARLTEL RRTSPGLWRFDLDSM\RNLK\AS LENSL\REVEAPYAL\QMEQL\N GILLHLESQLGQTPRTEAQRQA Q\EYEALL\NIKVKL\EAEI\ATLP GRLLGRMAKDFNL\GDS\LDER NSFQNL\QKTNTPPGNVDGKVV
18621	48989	A	18730	1	897	S\ETN\DTKVLRH MGHHRRQSLSPVLSLYSPDSMS FTTRSTFFTNYQSLGSVQAPSY GARAVSSAASVYAGAGGSGSRI SVSRSTSFRGGMGPGGLAAGM AGGLAGMGGIQNKKTMQSLN DRLASYLDRVRSLETENRRLES KIREHLENKGPQVKDWSHYCK TIEDLRAQIFANTVDNACIVLHI DNARLAADDFRVKYETQLAMR QSVENDIHGLRKLETEIEALREE LLFMKNHEEEVKGLQAQIASFR LTVEVDAPKSQDLAKIMADIRA QYDDLAGKNREELDKYWPQQF EENTTVVT\QSAEVGAA
18622	48990	A	18731	3	423	SLSPVLSLSPDSMSFTTRSTFST NYRSLGSVQAPSYGARPVSSAA SVYAGAGGLATGIAGGLAGMG GIQNEKETMQSLNDRLASYLDR VRSLETENRRLESKIREHLEKK GPQVRDWSHYFKIIEDLRAQIF ANTV\DNAR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18623	48991	Α	18732	663	2089	AFTTRSTFSTNYRSLG\SVQAPS
						YGAGPGRAARPASYAGAGGSG
						SRISVSRSTSFRG\GMGSGGLGQ
						RELAGG\LAG\MGGIQ\NEKETM
1						A/QALNDRLGLFTWDRVRSLG
						DRRTRKLGRAKFREHFGRKKG
						PQ\VR\DWSHYFKI\IEDL\RAQIF
]		RKYLLDNAPHPFLQDLTIAPSLL
						LD*F*EFKY*GQSLAMRQFLWR
						NDIPLGSAKV\IDDTQYHTDLQ
						AWRQRFEAPQRRELALSMKKE
						PRRRKLKGLSSPRLPSSWG*PVE
						VRLPPNFQDLRQRSLADIPGPN
						MNELGSKKNPRRKLRQSTWSQ
						QI*GRAPQLVTTTSLLEVGSLLK
						TTL/TQKLRRTSPGPWKIRPWTS
						MKKS*KAQLGRNSP*REVGRPP
		İ				YGPYKMEASFKRGSLLHP*SQS
						W/QTRPGAEGTAPQAPRSYEGP
						C*NIKRSKLEA*DRPPYPPALLE
						D\GEDF*SLVDPLDSKQLHAKPI
						QKTTTPPG*VGLGKVVS
18624	48992	Α	18733	53	1100	NFRVEAGVRGVQQKETCAFKV
						LESI\GKF\GLALAVAGGVVNSA
	:					LYNVDAG\HRAVIFDRFRGVQ\
	,	1				DIVV\GKGTHF\LIP\WVQETQLS
						FD\CRSRPRN\VPSQSTG*AKDL
						QNVQHHTAASLFRACRQPSFPR
						IFTSIGEDYGWSVCLPS\ITTEV\L
		İ				KVSVGSLWMLGELITKRE\LVS
						RQV\SDDL\TERAATFGLILDDV
						SLTHLTFGK\DFT*AVEAKQVA
						QQEAEKGQICWLEKAEQQKK\
						AAIISAEGDSKAAELIANSLATA
						GDGLIELRKL/E*AAEDIAYQLS
						RSRNITYLPTG\QSVLLQ\LPQLR
						GPPLPCTSRGLDLGPQPRLILNN
<u></u>		<u> </u>				SLSFLLPTPEINCEIFMIGLK
18625	48993	С	18734	222	458	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18626	48994	Α	18735	2104	2758	QFCPSRQKGCLQKTFRHFVAGH
						GGDFLFRSPTHAFPFCVPLFPLP
		ŀ				P/VTIVRPCPPFS*VLSSSPRNSQ
						NTIPRSPFLSPGTLSRSPAGSRDP
						GNPAPGSNTRLGPTAPVSRLAP
						VDIGFRSTDCWSLQPQVPVNPG
						SRPAPTNPRSRSTPVDPGSNTRL
						GPTAPVSRLAPV\V*ALGPLTAG
						PYSPR/CPVNPGSRPAPTNPRSR
			!			STPVDPGSNTRLGPTAPVSRLA
						PVDIGFRSTDCWSLQPQVPVNP
						GSRPAPTNPRSRSTPVDPGTRTT
						PEGL
18627	48995	Ā	18736	2	280	AMGFIVAPSLGCF/VGSRFVHG
						EGLRWYAGLQKPSWHPPHWV
						LGPVWGTLYSAMGYGSYLVW
						KELGGFTGVSRASEVAPPEPPP
1	İ					GSSVLCFLHA
18628	48996	Α	18737	331	972	TAAAAAMAPPWVPAMGFTLAP
						SLGCFVGSRFVHGEGLRWYAG
						LQKPSWHPPHWVLGPVWGTLY
						SAMG*VGVHWPGDKPGPLQGE
						AWPRTEDTGQLALNWAWPPIF
						FGARQMGGALVNLLLVS\GAA
						GHTTLGGRGAS\QWLAARLLYP
						\YLGLAWPSTTTTQTTCVWR\D\
						NHGWHGGRRLARVSARPTMG
			!			LQLHQAGAITLVMWWPSRFHD
						HWAC
18629	48997	Α	18738	1	603	
18630	48998	Α	18739	1	1782	
18631	48999	Α	18740	237	876	CESFCWHHFFWDIFNLFVTAPF
ł						LVHVDYQGFESVAKKYDVMN
ł						DMMSLGIHRVWEGFSCSGKMH
						PLPGD\QLLDVAGGTGDIAFRFL
						NYVQSQHQRKQKRQLRAQQN
ĺ						LSWEEIAKEYQNEEDSLGGSRV
						VVCDINKEMLKVGKQKALAQG
						YRAAHKSGVILDCLWFPPILHPI
						AQQICQLYLQNTSGIWPSLPYL
						CYHLLVQATIPSYLDCCNSL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	li .	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18632	49000	Α	18741	112	995	VLGSLSQEKRAAETHFGFETVS
						EEEKGGKVYQVFESVAKKYDV
					}	MNDMMSLGIHRVWKDLLLWK
						MHPLPGTQLLDVAGGTGDIAFR
						FLNYVSVPASEKTEEAVKGPTK
,						FILGKNLPKSTRMKKIPWAG/TR
		1				VVVCDINKEMLKVGKQKALA\
:						KDTRAGLAWLL*DAEELPFDD
		1				DKFDIYTIAFGIRNVTHIDQALQ
						GSSFGCLKPG\GRFLCLGFSQVE
						QSP\LSRLYDLYS\SRVIPVLGEV
			ľ			IAGDWKSYQYLVE\SIRRFPSQE
						\KFKD\MIEDAGFPQ/V*LTKV*H
						QGIVAIHSGFKL
18633	49001	Α	18742	117	403	
18634	49002	C	18743	62	370	
18635	49003	Α	18744	179	527	IQEPPVI**GLHMVHNFNTIFVSS
						YFKFESTFAISFLKVFNT/CFFSI*
						IKSLDEQCVVGKISKHWTGILR
						EAFTDADNFGIQFPLDLDVKMK
						AVMKVGICGKNLVKQGNFTMT
	<u> </u>	ļ				HSN
18636	49004	A	18745	3	1242	AAPQAGLSPVAIAAAIQLHLHS
						TQCSSPNTCCLPRRTRATIYYSR
						WSYHPLGSVP*SP*PFQEAS/AL
						TLPPACSFYGPLT*FQPKP*GSFP
						LSQ*MEYTIGLYT*TFHCPGTSR
						RQIPSSYLNCKDAFLPLL/SNPP
						QCRPFTGVGLVDVLTGFETNN KYEIKNSFGQRVYFAAEDTDCC
						TRNCCGPSRPFTLRIIDNMGQE
						VITLERPLRCSSCCCPCCLQEIKS
						LDEQCVVGKISKYWTGILREAF
		}				TDADNFGIQFPLDLDVKMKAV
		1]		MIGACFLIDRNCSPAMEQSWM
						ENYFDEMTEIGFRRSVITNFSEL
		1]			KEHVLTHCKEANKNLDKMLDE
						WLTRKNSVEKTLNELMEVKTI
		1				NEKLTIGKISKYWSGFVNDVFT
		1				NADNFGIHVPADLDVTVKAAM
		1				IGACFLFAFRLGSELHN
18637	49005	В	18746	277	2415	10.101 B. M. REGOEDIN
18638	49006	A	18747	102	196	
10000	1,,,,,,	14.	110171	1.72	1170	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
18639	49007	A	18748	897	1656	SNAGLRHVCRTSCQSGRARSCH
						VGLHTPWVS*VPWHPWGIRWR
	1					GDPQKSQRSRKCCVPGGAP/QK
						GGDPEPQPSLSVSWNWLSGSCA
						GFRAVLPRKPQRKVWLFASSA
						GTAATEPPR**LAWKADSGGDP
		İ				WQQPQQSARPHA*TWTWRGD
		į				GAPPGRVA/QGRGTAGGASAG
						VSSGRSRGSPQTGFQALQSCLE
						ASRRPPVVACNPHPAGALPAAQ
						AGNRATLMTGSCLQMPIMYEE
						AQLRLEELCTKARLNCQWFL
18640	49008	Α	18749	756	884	CLQRLLPPGFRRQPSKYRRNRG
l.						KSGKG*RPSPVATWQPDYNI
18641	49009	Α	18750	1826	2772	RRVLEAMGPSRAAKVLSVCVR
İ						SRIGPLICSQRHPATSKTKPQGR
						PPGEQGTPRDNPRRNNRPARSA
						TPRGA/PQGAPPRPGEEPNAPQT
	1					IDKGPPSPPSRGHTRRLARSGNQ
						PPERHTPAAGGKTRYVRQSVL
						HRARGGEPVRMPRLPRIAEA/P
	[APDRVECF*VVQSPAREMHSCQ
						ADSMCPVW*GRIWRSAGSC*SR
						SEKILVWIPQSGLQAHACPSEA
						HKMPGNPRDGGAGICKSSPGG
						KGERRKVEGGSTTRAGTSRDS
						VSKRHANTMGEQAQGEKHPRG
						TV*KTKRRGPKGQTRGTKAPEE
						ETSREPPCTGPI
18642	49010	Α	18751	196	322	
18643	49011	С	18752	189	650	
18644	49012	Α	18753	1	53	
18645	49013	A	18754	31	449	
18646	49014	Α	18755	1	389	GMPTSTIASCSQDGRVFIWTCD
						DASSNTWSPKLLHKFNDVVWH
			1			VSWSITANILAVSGGDNKVTEG
						QQNEQ*QDRWGLAPHPPAPGL
						PLPGPTNQTTGKSPQLQQDYFP
		<u>L</u> .				RRSYRCSHRLIICLNVIGNAL
18647	49015	Α	18756	1	1041	
18648	49016	В	18757	15	782	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18649	49017	IA	18758	1	1425	EVKKINNAHTIGCNAVSWAPA
		1		_		VVPGSLIDHPSGQKPNYIKRFAS
	:					GGCDNLIKLWKEEEDGQWKEE
						QKLEAHSDWVRDVAWAPSIG/
	Į.					LWPPSTIASCSQDGRVF\IWTL**
						CPQAITVVPLNCLHKFNDVVLG
	Ì					MLNWGPSQPTSLAVSGGDNKN
		ŀ				GTNTVCPTGWLQDSNTLVHLV
		1				TYKMPRSLGLGFLWCQLLSPR
						KYREGTSPVSLWVGEVWFINP
		İ				DWWGSIRVYHPSPSGSYRTHKP
						PLNNLSCDTSIIRELLKGKKFLL
		l				FVSVYPVPSTVLGRKKVSVINT
						VDTSHEDMIHDAQMDYYGTRL
						ATCSSDRSVKIFDVRNGGQILIA
	1					DLRGWPRASRYQQGHQDLFIL
						RSDLPSQVFIRDKLMERRNRRT
	1					GRTEKARIWEVTDRTVRTWIGE
						AVAAAAADGVTFSVPVTPHTF
					ļ	RHSYAMHMLYAGIPLKVLQSL
į						MGHKSISSTEVYTKVFALDVAA
						RHRVQFAMPESDAVAMLKQLS
18650	49018	В	18759	323	1583	
18651	49019	Α	18760	3	419	PSASSSPARIPAATRQGRRSPRIP
						PAPSNEAPRLPADREFLEQQPPR
		1				FPKPTIQASGADRAVDCGILKL
						QKS\PARLARPRWPRRPSKRFW
						SAAGSVEEQPKPPRRRAKSPEQ
						SPQLSVLPLVTITPQASGASFEK
						MKN
18652	49020	Α	18761	2	385	WGRGGGRGLLYGRRHFVRRRR
		1	1			RGGRSPGESPALTVT*GAV\PSA
		1	}			AFRESAQPGPRSSEAHISTSPKR
		1				TDGGRDS*PGLPRSSLTPSLPRP
		1	1	<u> </u>		ASFSTSPGRVSAPHRLPRHS/AG
						PGLSGSS*PQPYSDPT
18653	49021	Α	18762	242	422	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18654	49022	Α	18763	2	992	AREKKQKTRRQNKNIPRQINKT
						QRICC*QSCLTMTQDFFWPLCQ
					1	GHSRGAPCLLVPPGLSLTCAPA
						QPAARPGVSQLTCAIACTHIQW
					-	FPSSCPPSEKNGDTLTV\KTPSA
						SSSPARIPAATRQGRRSPRIPPAP
						SNEASPAPGRPRVPGAA\PPRFP
						KPTIQASGADRAVDCGILKLQK
					,	S\PARLARPRWPRRPSKRFWSA
l						AGSVEEQPKPPKAP/PAKSPEQS
						PQLSVLPLATITPQAYGASFEK
						MKNLKTRRGGRSRWHFPFNVIP
		1				QAPSLKWGFQTSDASSAEVFH
		1				VDRSLASHKMGPGRRPRSPLW
						APPFCEAAAARRQEPGVCIMVE
18655	49023	A	18764	292	348	
18656	49024	A	18765	2	466	LITVVKLLDTMVDTFLQKLVA
1.0050	.,,,,,		1.0.02			AGSYQRFTDCYKCFYQLQPAM
						TORIYDKFIAQLQTSIREEISDIK
						EEGNLEAVLNALDKIVEEGKVR
						KEPA/WQRDTLRRHVQKQEAE
						NQQLADAVLAGRRQVEELQLQ
	1					VQAQQQAWQALHREQRELVA
			}			VLREPE
18657	49025	Α	18766	2	511	FNMAEASSANLGSGCEEKRHE
		}				GSSSESVPPGTTISRVKLLDTMV
						DTFLQKLVAAGS/YSTPSLGRK
						CWS\RLPSGGQSVKQAFSWAAC
				1		RLPQGRKEEISDIKEEGNLEAVL
						NALDKIVEEGKVRKEP\AWRPS
						GIPEKDLHSVMAPYFLQQRDTL
						RCHVQKQEAENQQLADA
18658	49026	Α	18767	224	748	NRPSVGRRAGYHRGGSYQRFT
						DCYKCFYQLQPA\MTQRIYD\K
						FIAQ\LQTSIREEILFDNQKQEGE
						PRKLSWNALG*KFVGTKGKVR
						KEPAWRPSGIP\EKDLHSVMAP
						YFLQQRDTLRRH\VQKQEAENQ
						QLADAVLAGRK\QVEELQLQV
						QAQQQAWQALHREQRELVAV
						LREPE
18659	49027	Α	18768	1	421	l.,,

SEQ ID	1		SEQ ID NO:	P .		Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
18660	49028	A	18769	666	1445	AGITTIEAAKRTIQVLHLQAND
		l				AQERA\ERSQREFREERRAREPG
						LRLK*SFLNRRIQ\LVEEEL\DRA
						QERLGHLPCKKL*RS*KKLADE
						SERRY*RLFEN\RALKRLKEKIG
		ĺ				TSREIPTQKKLKHICRKEAD\RK
						YEEV\ARKLGDHLKEDLEPTE\E
						RA\ELAESRC*EIDEQIRLIDQNL
						QCLSAAEDKYSQKEDKW\EEE
						MKI\LTNKLREAET\RAEFAERS
						VAKLKKTIDDLEDKLKCTKEEH
	1					LCTQRMLDQTLLDLNEM
18661	49029	Α	18770	2	158	SAAPGAVPPPEADSTSAGMSRR
						PCSCALRPPRCSCSASPSAVTAA
						GRPRPSDSCKEESSTLSVKMKC
						DFNCNHVHSGLKLVKPDDIGRL
						\VSYNP\AYLEGSCKDCIKDYER
						LSCIGSPIVSPRIVQLETESKRLH
						NKENQHVQQTLNSTNEIEALET
	1					SRLYEDSGYSSFSLQSGLSEHEE
	1					GSLLEENFGDSLQSCLLQIQSPD
		l				QYPNKNLLPVLHFEKVVCSTLK
						KNAKRNPKVDREM\LKEIIARG
	1	İ				NFRLQNIIGRKMGLECVDILSEL
						FRRGLRHVLATILAQLSDMDLI
	,					NVSKVSTTW\KKILEDDKGAFQ
						LYSKAIQRVTENNNKFSP\HAST
						REYVMFRTPLASVQKSAAQTSL
						KKDAQTKLSNQGDQKGSTYSR
ı						HNEFSEVAKTLKKNESLKACIR
I						CNSPAKYDCYLQRATCKREGC
l						GFDYCTKCLCNYHTTKDCSDG
ı						KLLKASCK\IGPLPGTKKSKKNL
						*RLHVGWHEPAPLQLRPTATPL
1	1					LLQRQPQRSDSRRAPSTLG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence	ŀ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18662	49030	A	18771	4	1276	RKAASSYVSRMALPIIVKWGG
						QEYSVTTLSEDDTVLDLKQFLK
l		ŀ				TLTGVLPERQKLLGLKVKGKP
						AENDVKLGALKLKPNTKIMMM
						GTREESLEDVLGPPPDNDDVVN
}	1					DFDIEDE/R*FEVENREENLLKIS
		l				RRVKEYKVEILNPPREGKKLLV
						LDVDYTLFDHRSCAETGVELM
	İ			ļ		RPYLHEFLTSAYEDYDIVIWSA
						TNMKWIEAKMKVKPLGVIWG
						KFSEFYSKKNTIMFDDIGRNFL
		i				MNPQNGLKIRPFMKAHLNRDK
						DKELLKLTQYLKEIAKLDDFLD
						LNHKYWERPRVRCHIFHGKED
						RYDPCPMGACSCKTVCTTNLA
		į				LPALPLGSMKNGQPCRNVIGQK
						GYALTVINSGPQQGLFVRILLG
į			1			LFIRDFFPAGCGARPFWNEGLM
Ì						AYFQRRTLVGVMLVLSSSKSFK
18663	49031	C	18772	113	257	
18664	49032	Α	18773	143	330	HHGPVPFPRGLIS*HSLHCPCPT
						PGGKNVFFFCEYFLKLAGIETT
						NCSADKRGKKVCCLIPR
18665	49033	Α	18774	3	432	
18666	49034	С	18775	112	192	
18667	49035	Α	18776	3	1015	SSRPVRPRPAARLSAMSSTQFN
						KGPSYGL\SAQ\VKNRLL\SKYD
		1				PQKE\AELRTW\IEGLTGLSIGPD
						FQKG\LKDG\TIL\CTLMNKLQP
						G\SVPKINRSMQN\WHQLENLS
						NFIK\AMVSYGMNP\VDLFEAN
						DLF\ESGNMT\QVQVSLLALGG
						KRPKTKGAAEGGLDIGVKYSE
						KQERNFDDATMKAGQCVIG\L
İ						QMGT\NKCASQSGMTAYGTRR
				1		HLYDPKNHILPPMDHSTISLQM
						GT\NKCASQVG\MTA\PGTRRHI
						YEYQAGNPTSCDNFSM\SLQ\M
						GYTQGAQTQSGQVF\GPGRPDI
				1		*TPSTCPQGTI\ADGAPSGTGDC
		L				PDPGEVPEYPPYYQEEAGY
18668	49036	A	18777	2	110	
18669	49037	Α	18778	74	289	

SEQ ID	SEQ ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18670	49038	Α	18779	138	2039	GAAMAAAGRLPSSWALFSPLL
10070		` `	10715	1.55		AGVALLGVGPVPARALHNVTA
				1		ELFGAEAWGTLAAFGDLNSDK
						QTDLFV/LAGKK*LNRLFGRPE
						CTRILNPK*RYLSRITVH**QV*S
						LGAYDGDSQMDVLLTYLPKNY
						AK\MN*ELLSSGDKIKH*ILTI*P
						YSIGL/LQDEPLIMDFNGDLIPDI
						FGITNESNQPQILLGGD/CYHGI
						QH*PLQVKCEFHI/PHAFIDLTE
						DFTADLFGATLNGTTSTFQFEI
						WENLDGNFSVSTILEKPQNMM
						VVGQSAFADFDGDGHMGSFTC
						QGCGR*K/CAKRVPSYLVRSGM
						KQWVPVLQDFSNKGTLWGFVP
						FVDEQQPTEIPIPITLHIGDYNM
						DGYPDALVILKNTSGSNQQAFL
						LENVPCNNASCEEARRMFKVY
						WGADRPKSN/SKGAMVATFFDI
						YEDGILDIVVLSKGYTKNDFAI
						HTLKNNFEADAYFVKVIVLSGL
						S\SNDCPRKITPFGVNQPGTLYH
						VYNCRCKWVSEKWIRWPNSAQ
						SAHLALQLPYNVLGFRVGSANF
		ļ				LDHL\SLGIPRPSWRKIFYGKQE
						WTAIIPNSQLI\VIPYPHNVPPKL
						GVPNCILYTKLILFWLTAIALIR
						CLCFQSLAINWHFTLGREKKAG
						WIEEKRTRKAHRFPFWMAMW
18671	49039	A	18780	1	966	WIELKCIKCAIRCIT WMANW
18672	49040	A	18781	1	1035	RPPFPVPGVQKCPLPTRGGLHIG
				<u>.</u>		RWLRDRAGPPEAQDGTGGRSR
						SRRRPPALPNSRPSPVASGREM
		ļ				VVLSVPAEVT\VILLDI\EGTTT\P
						IAFVKG/DILFPYIERKMLK\EYL
						QTHW\EEEEFQ\QDVSLFEGNK
						A\EEDAHLDG\AVPIPGKHLGIG
		İ				V\DD\LQQMI\QAVVDNVCWQM
1		ļ				SLDRKTTALK\QLQGH\MWRAA
1						FTAGRMK\AEFFADVVPA\VR\K
						WREAG\MKVYIYSSGSVEAQKL
						LFGHSTEGDILELV\DGHF\DTKI
						GHKRRRVKVTRK\IADSIGCST\
						NKHFVFWTDVTSRRASAGLRE
Ì						AGCCTLAVV\VRPG\NAGINR*L
						RKTYYSLITSFQWNYYLPSST
L	L	L	L	L		KKI I I DELIDI (WINI I LEI DO)

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18673	49041	A	18782	11	528	DNELLDYEDDEVETAAGGSMS/
						EAPAKKDVKGSYFSIHSSGFRD
						FLLKPELLRAIVDCGFEH\PSEV\
						QHECIPQAILGMDFLCQAKSGM
						GKTAVFVLATLQQLEPVTGQV
						SVLVMCHTRELAFQTR*KTRPF
						WNVTVCPSGEDTRVGVKETLL
						PPPLTAPTPWLPSFASPPLLNPH
18674	49042	Α	18783	207	309	
18675	49043	Α	18784	85	348	PPHSALLPFPTPFLPPSQLEIFVD
			,			DETKLTLHGLQQYYVKLKDNE
		İ				KNRKLFDLLDVLEFNQVS*TSS
						RGMSIGALQL*QKPGY*VHFY
18676	49044	Α	18785	1	930	CCCRHTRSPCLVMAENDVDNE
						LLDYEDDEVETAAGGDGAEAP
		1				AKKDVKGSYV\SIHSSGFRDFLL
						KPELLRAIVDCGFEHPSEVQHE
						CIPQAILGM\DVLCQAKSGSGER
						QAVF\VLATLQHLEPV\TGQGV
		:				CAG*CCHTRELAFQISKEYERFS
						KYMPNVKVAVFF\GGLSIKKDE
						EVLKKNCPAYPSVGTPGRILA\L
						ARNKSLNLKHIKHFILDECDKM
						L\EQLDM\RRDVQE/IFFRMTPH
						EKQVM\MFSATLSKEIRPVCRK
						FMQDVNTLLPSLPPLPARCLLPF
						LALFLRLPCHSSAKKAACAQLG
18677	49045	Α	18786	96	1618	LFTPCFHLFCENPSRSPFPSSPAG
						PVMAENDVDNE\LLDYER*MR
	ļ					WETAAGGDGAEAPAK\KDVKG
		İ				SYVS/LSHSSGFRDFL\LKPELLR
						AI\VD\CGFEHPSEVQHECIPQAI
						LG\M\DVLCQGKSG\MGKTAVF
						VLATLQQLEPVTGQVSVLVMC
						HTRELAFQISKE\YERFSKYMPN
		1		}		VKVAVFF\GGLSIKKDEEVLK/R
1						RNCPHI\VVG\TPGRILS/LWVRN
						KSLNLKHIKHFILDECDKMLEQ
						LDMRRDVQEIFRMTPHEKQV\
1						MMFQCYLGAKEIRPVCRKF\M
						QDPMEIFR/VDDETKLTLHGLQ
1						Q\YY\VKLKDN\EKN\RKLFDLL
]					1	DVLEFN\QVVIFVKSVQ\RCIAL
						APAY*WEQNFPA\IAI\HRGDAP
]			RKERLS\R*SSSFKDFPTNEILIG
						YPTLFGRRHGTSEAGETIAF*FM
						NMP*GILKTLPCHR\VGPEQGRF
						GHPRGL\AITFVSDEE*LPRSCD
						DVQGSALRVKFSELPDEIGHLP
10670	10046	 -	10707	1,	262	PTLEQDTGRRLAHFGNVDRLFL
18678	49046	Α	18787]1	363	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
18679	49047	Α	18788	78	714	GTTEGKRQEAVGYLCVLPLTTL
		İ				PLVTRLPPQVADTMLPPMALPS
						VSWMLLSCLILLCQVQG\EETQ
						KELPSPRISCPKGSKAYGSPCYA
					:	LFLSPKSWN\DADLAC\QKRPSE
						KLVSVLSGAEGSFASLLA\RSIS
						NSYSYIWIGL\HDPKKG\SEPDG
						DG*EWSSTDVMNYFAWEKNPS
						TILNPGHCGSLSRSTGFLKWKD
						YNCDAK\LPYVCKFKD
18680	49048	Α	18789	52	186	QSEVHHGIISYTTGQGENLSEPI
						WKSWQRSQDL*QENHPEIYSLT
18681	49049	A	18790	666	2732	KQLIKMCYFILLLILSSHIGKD
						KAIDKNGILVQYWQE*KLVEPF
						*KAIWPPCMETLNVDTPFPQPV
						CLGIYPNE*KLSK/CFKP*APKV
						YFHGCL*Y*NHWEPTETIPPVK
						GWIK*ILF
18682	49050	Α	18791	112	340	
18683	49051	Α	18792	3	323	LSSLASMSFTTCSAFTNYWSPG
						SVQVPSYGTQPVSHAASVYAG
						LGGSGSRISVSH\SPMAGGLAG
						MGGIQNEKETMQSLRDRLASY
10604	100.50	ļ. —	10502	,	004	LDRVRGLETENWKLESKIQEH
18684	49052	A	18793	1	994	APSYRA/RLVRSVASVYAGPGS
						SGSRISMSRSTSFQGGLGSRSM
			:			AA/GMAGGLAGMGGIQNEKET
						MQSLNDRLASYLDRVRSLEMG NWKLESKIWEHLEKKGPQVRD
			1			WGHYFKTIEEDLTQIFTSTVDN
						TCIILQIDNAHLAADDFRVKYE
						TEPATCQSVENDIHGLHKVIDD
			i			TSVTQLQLETEIEAL\KEALLFL\
		[KKNHEEES/WRHTVQALEIDLD
						WMKNLEASLENSLK/EGVEAC
						YTLQMEQLNGILLHLESELAQT
						WAEGQGQAQEYQALPNIKVKL
						EAEIATYRRLLEDGEDFNFGDA
			l			LDSSNSTQTIQKTPTC/RTVDGK
						VVSETNDTKVL
18685	49053	A	18794	3	377	
18686	49054	В	18795	54	136	
18687	49055	A	18796	149	419	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ļ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18688	49056	Α	18797	83	1402	AEKAGGKKGEAPSYRA/RLVRS
						VASVYAGPGSSGSRISMSRSTSF
						QGGLGSRSMAA/GMAGGLAG
l						MGGIQNEKETMQSLNDRLASY
İ						LDRVRSLEMGNWKLESKIWEH
ĺ						LEKKGPQVRDWGHYFKTIEED
İ						LTQIFTSTVDNTCIILQIDNAHL
						AADDFRVKYETEPATCQSVEN
						DIHGLHKVIDDTSVTQLQLETEI
						EAL\KEALLFL\KKNHGEEIKGL
						SRRQIASSGVDPWRVDAPKSSG
						PSPKIMGRHPGPKYDEL\ARKR
						TREE\LAKYSSQ\QIEEST/TQLV
[TNTSLL/GGLELAETNLHRS*TY
						KSRSLEDSTLDSIEKS*KAQAW
İ						KKQPLKGSWKAPLRPLPDWSK
						LQTGILACTFESRAGNKTRARG
						TSAPGPQEL*GPLLKHQKVKAG
						GLRSPTLPARLLEDAEAL*SLV
						DPLDSKQLPCNTIQKTTTPPG*V
18689	49057	С	18798	140	283	
18690	49058	Α	18799	2	783	VVQFARREKPSSSKVSQRHPCT
		l				EAVREVLHG\NQRKSPQRSWET
		}				VELQISLKNYDPQEGQSAFSGT
		l				RQALSPLPRP\KFSCVCPGGTQQ
						H\CDEA\KARGISPPHGTFEGA*
		l				KNLNKE*KNWSKKAGPRKY*L
		1				RFFGPQKSLDQSKIPPNPSGPGF
		1				K*RAGK\FPFPWFTHN\ENMGG
		1				PKLDEGEVPPIQVFQMKERCLC
ĺ		1				L\AVAV\GHV\KMTD\DELVYNI
		1				HLGCQLSWVSLLKEKNW\QNV\
		<u> </u>				RALYIKSTHGQSPKRLIFKAHFE
18691	49059	С	18800	239	555	

SEQ ID	-	1	SEQ ID NO:		l .	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
18692	49060	I A	18801	401	3014	TRASSTMSKSFQQSSLSRDSQG
10092	75000		10001	401	3014	HGRDLSAAGIGLLAAATQSLSM
						PASLGRMNQGTARLASLMNLG
						MSSLLNQQGAHSALSSASTSSH
						NSQSIFNIGSRGPLPLSSQHRGD
						ADQASNMLASFGLSARDSDELS
						RYPSDTITPENLRPILLQLKRRR
						TEEGPTLSYGRDGRSATREPPY
						RVPRDDWEEKRHFRRDSFDDR
						GPSLNPVLDYDHGSRSQESGYY DRMDYEDDRLRDGERCRDDSF
						FGETSHNYHKFDSEYERMGRG
						PGPLQERSLFEKKRGAPPSSNIE
						`
						DFHGLLPKGYPHLCSICDLPVH
						SNKEWSQHINGASHSRRCQLLL
						EIYPEWNPDNDTGHTMGDPFM
						LQQSTNPAPGILGPPPPSFHLGG
						PAVGPRGNLGAGNGNLQGPRH
						MQKGRVETSR\VV\HIMGFSTE
						GKTLRYQLLQLVEPFGVHFKFI
						*FLNKNLLRAFIEMATTE\DCFK
						AAV\DYYTTTPA\LVF\GKPVRV
						HLSQKYKRIKKPEGKPDQKFDQ
						KQELGRVIHLSNLPHSGYSDSA
						VLKLAEPYGKIKNYILMRMKS
						QAFIEMETREDAMAMVD\HCL
						KKALWFQGRCVKVDLSEKYK
						KLVLRIPNRGIDLLKKDKSRKR
						SYSPDGKESPSDKKSKTDGSQE
						AERST\EGKEQEEKSGEDGEKD
						TKDDQTEQEPNMLLESEDELLV
						DEEEAAALL\ESGQFSGETSTDL
18693	49061	Α	18802	3	1013	NCRAAWRAKLEGCLSLRRNAC
						QCFLIWTQTPDREQSYSLDQLP
						REVLNQEEKSISIWDIYLHLEM
						GNGLSDQTSILSNLPSFQSFHIVI
						\LGLDCAWKRQLSYTRLQFQLN
						L*NT\VPYPKGFNTEKI*/RVTLG
	1	İ				NSKTV\TFHF\WDVGGQEKLRP
	<u>}</u>					LWKSYTRCTDGIVFVVDSVDV
						ERMEEAKTELHKITRISANRGV
		İ				PVLIVAN\KQDLRNSLSLS\EIEK
						LLAMGELSSSTPWHLQPTCAIIG
						DGLKEGLEKL\HDMI\IKRRKML
						RPTRKRKDEYQFPNIICVGVGF
						LWSDFDKNRRVSTAGVCLSAPP
						WMLLKLCFVEQFRCPTRVALW
						KMSKCSAS
18694	49062	Α	18803	3	1103	
18695	49063	В	18804	34	734	
	49064	Α	18805	1	1741	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			: -	sequence		
18697	49065	ĪA	18806	6460	6635	RTSTERSHLKTRYQRTSCSSNR*
	.,					LNSSSSKTSFRCSLTTSLFWRHT
						IFNMLLINRRLL
18698	49066	A	18807	5567	5853	IGILEA*LLRNLPNDSIKALVDA
10050	1,7000	'	10001			*YKSLLSAVERCSFMVLFGEPM
		1				NKPPLAILRIQSGVNRPPVISIM
		}				ALIGPLVTPSASTYSGVTNADLS
		1				SATK
18699	49067	A	18808	664	992	SCADIHGYHRGNSSNNDSNLIP
10055	1,5007	ľ.	10000		,,,,	KQTNRDCLNNGHQNHSCNG/D
						*PQPQR**QLTLQNLMQHQ*KQ
		1				QYPLPQHQQQE*RTR
18700	49068	l _A	18809	3	1698	Q 11 El Quiqqq E KIK
18701	49069	В	18810	227	883	
18702	49070	A	18811	180	373	PDTLRSILPMLLEQLFCRTFGLL
						SP**RDTSLHLGHKTEQHT*LV
İ		1				MGSWFYLLASQQPRCCYAC
18703	49071	Α	18812	495	701	
18704	49072	Α	18813	138	418	AYRLLLCLVYSEGFYMHLEFQ
		1				NRLYHQLVF*GRLWHFW\KSIK
		1				RFSIY*FFWYRW*YICKLNKGR
		1				LQDVIGVINT*CHFLS*GLH*TIP
		1				GIHRL
18705	49073	С	18814	62	175	
18706	49074	Α	18815	644	828	
18707	49075	Λ	18816	1	1107	
18708	49076	В	18817	84	342	
18709	49077	A	18818	3	1399	
18710	49078	В	18819	1	1377	
18711	49079	Α	18820	1	546	
18712	49080	Α	18821	855	1099	VTLYSEYQHPSVREYGLYLTSH
						RIEPLECVHRTL*PVLLSMQPSV
						LEYYEQQGYEAEY*LQEECHH
10710	10001	 	10000	ļ.,	1.600	QRTSSL*DLLKQLLQ
18713	49081	A	18822	1	1680	
18714 18715	49082 49083	A	18823 18824	2	3951 6446	
18716	49083	A	18825	2497	21785	SSPYPFTKQSSTNNFVFDGNELF
18/10	49004	^	10023	2497	21763	NSRDRLSQRLFYILQSSVHLINR
						Q/LEGYVQHHLNLAEI*HHGLG
						` ` `
						KDAYIAAMYNNDDGATFQMEI
						GGIHTNAALAAGDKLQCIWRY
		1				LDVNGNAPAAENEFEVVSYEG
						FGTSNFSVKCKGMLPGKFIGCD
						AFYGKYLQTDGPIKQVDSVKW
		1				FTNLTVSGSGRKQLGQRKYPQ
						VVMGMGMTSGFDDGYNLTPE
		1				RQVKMAYGLGYRDWWTTYIG
L	10005	 	1000 (<u> </u>	170	MSHYWKGLTAFQDKETGELI
18717	49085	Α	18826	31	479	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	i e	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			<u> </u>	sequence		
18718	49086	A	18827	255	1002	SQFSLSQVLVDSAEEGSLAAAA
						ELAAQKREQRLRKFRELHLMR
					<u> </u>	NEARKLHHQEVVEEDKRLKLP
						ANWEAKKARLEWELKEEEKK
						KECAARGEDYEKVKLLEISAED
				İ		AERWERKKKRKNPDLGF\SDY
						AAAQLRQYHPV*PKQIKP\DME
						TY\ERLREKHG\EEFFPTSNSLLH
İ						G\THVAFHRGKFDRVVLRSGKN
1		ŀ				QFEKRD/KF*ARRRPYNDDADI
			1			D\YINERNAQFNKKAERFY\GK
						YTAEIKQNLERGTAV
18719	49087	С	18828	91	225	
18720	49088	Α	18829	3	212	
18721	49089	Α	18830	88	239	
18722	49090	Α	18831	865	1898	QQQQQQHRRYPRRRSPRAWC
						SHPRSPLLFLPSLAPWPDQLTEE
						QIAEFKEAFSLFDKDG\DGTITT
				l		KEL\GTVM\RSLGQEPNKKLEL\
				1		QDMINESGMLMG\NGTHLTFPR
						IF*LMMARKMKDTDSEEGN/LR
						EAFRV\FAKDGNGYISAAELR\H
						VMTNLGEKTTRF/KKVDEMIRE
						A\DIDGDGQVNYEEFVQMMTG
		1				KMEDLTFQLPFSPPLEESNWNL
		1				LLTFLQKKEKKKKVHLFHSCFL
						YSKTECQKYLLVHTHQNLHVL
				1		VGGPCPLKDQATHQFYNINTCT
						NLNDNGLLKRSILLMINTLFGL
						ASFFHACSFDDWSTSQGLWYL
						KLENGKNQISKPIPNGVLVHFV
18723	49091	Α	18832	1	330	WVVLKNCVGVGLLIAT*MWFI
						SNKYLVQRQSRDYDVEWGYAF
		1		1		DVHLNAFYPLLVILHFIQLFFIN
		1				PLPFLKNTVILLYPFAPLILLYGL
		<u>L</u>				SLALGWNFTHTLCSFYKYRVK
18724	49092	Α	18833	1	534	VPSNSVNSLVQGNGVLNSRDA
						ARH/TAGAKRYKYLRRLFRFRQ
						MDFEFAAWQMLYLFTSPQRVY
		1		1		RNFHYRKQTKDQWARDDPAFL
		1				VLLSIWLCVSTIGFGFVLDMGF
						FETIKLLLWVVLIDCVGVGLLIA
						TLM/WHCHFWKKQVILLYPFAP
				1		LILLYGLSLALGWNFTHTLCSF
						YKYRVK
18725	49093	С	18834	295	570	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	T .	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	li .	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
18726	49094	A	18835	418	1567	I SFTLKNCVLQRVSSILPSNTAEV
10,20	1,505,	ľ.	10055			DFPSWTTQSDWQQTRAQVLLQ
						ARARAALRLVLIILRHDGASPP
1						MRTRLSPTATLATGSGASEAPV
,				-		SGDRKPAATTSGGPRRKMLPST
						SVNSLVQGNGVLNSRDAARHT
				İ		AGAKRYKYLR\RLFRFRQMDFE
		1				FAAWQ\MLYLL\TSP*RVYRNF
						H\YRKQTKDQWAR\DDPAFLGF
						PFKYSGFCVSTIGFG\FVLDMG\
						FFETIKLLLWVGLID\CVGVGLL
						\IATL\MW\FISNKYLVKRQS\RD
						YDVEWGYAF\DVHLNAFYPLL
1						VILHFIQLFFINHVNLTDTF\IGY
						LVGNTL\WL\VAVGYYILR*LSL
						GYSVGLLFFSSIAHF*KNT\VILL
						YPFA/PL*FLPPTGFPLALGWNF
						THTLCSFYKYRVK
18727	49095	Α	18836	1	888	MPGLLLLRGSQNLKQQAEILCG
						LGLALVISAVKPLSKKFMLPNP
						TSMKPESDGIYSINLKRTWEKL
						LLAGHAIVVIENSADVSVISSRN
						TGQKAMLKFAAATGATPIAGH
						FTPGTF/TNQIQAAYWEPRLLV
						VSDPRADHQSLTEASCVNPPAI
						ALCNTDSPLCH/VDIAITCNNKG
						APSVGQMW*MLAWEVLHMRG
					,	TISCN/SPWEVMSNLYFYRDPEE TEKEEQAAAEKAVTKKEFQGK
						WTVSAPEFTVTQPEVADWSEG
						MQICNLGKAQQGQLSSDAEDD
				1		FWSLESLDGYSHTVTFR
18728	49096	A	18837	74	848	REIFINVRSLLMACFVKEEDVL
10720	7,070	1	10057	'	0.10	KFLAARNP/HVGGTNLDFQME
						QYIYIRKSDGIYIIRLKRTWEKL
						LLAARAIVAIENPADVSVISSRN
						TGQRAVLKFAAATGATPIAGRF
		ŀ				TPGTFTNQIQAAFRE\PRLLVVT\
						DPRADHQPLTEASYVNLPTIAL
						CNTDSPLRYVDIAIPCNNKGAH
						SVGLMWWMLAREVLRMRGTI
						SREHPWEVMPDLYFYRDPEEIE
						KEEQAAAEKAVTKEEFQGEWT
						APAPEFTATQPEVADWSEG
18729	49097	Α	18838	72	528	DEACGVCCYSPGASPMPP*GRT
						RKEGRTAGKVRPKERPKTPVN\
						KSGGQGPKRKNWSKRQKFRDK
	1					L\NN\LVLFDKA\TYDKLCKEVP
	1					N\YKLITPSLWSPERL\KIRGSLA
1						KGKPFQEAP*VKGLI/RNLVSKH
L	<u> </u>			l	<u> </u>	RAPSNFTPGNYQGWEDAPSLL

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO:	Nucleotide location of first	Nucleotide location of last	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
110.	sequence	nou	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
18730	49098	В	18839	6	125	
18731	49099	В	18840	249	291	
18732	49100	Α	18841	128	669	RPGHAQTLGSPSLALGMAGQF
						RSYVWDPLLILSQIVLMQTVYY
						GSLGLWLA\VVDGLLRISPSLD
						QMFDAEILGFS/TPFQGRLSMM
						SFILNALTCALGFAVISSGEEKQ
						CLDFTVTVHFFHLLGLLGSNSS
						PFSPSGG*TLVGLGPKPVVALH
						SLAVIGEYLCMRTELKEIPLNSA
18733	49101	Α	18842	1	818	MAELTVMRFHTEPQGSEGLRLP
						RISGSKPKQMHVTWLEPLENGI
						CNGNTGNAQLLIPPGVIFHKYP
						SHYVTAMLKSLKWPPLQTELTS
						SSGQALWGLVGGGSPHRQAST
						EKLFSEKVPKTAENFPCSELSCF
						HGENGFGL*GVPCFHRLFPRVY
		l				AVRGGDFHTAINGT\GGKSHPM
ŀ		İ				GKKFEG*ETSSLKHTG\PGDLVP
						WQNAGP\NTNGSPVFLTSCTAQ
		1				ELSGLDGKPCGVLAKVKE\GM
1		1				NIVEAMER\FGSR\NGKTSKKITI
						A\DCGTTPNKV
18734	49102	Α	18843	4	454	
18735	49103	Α	18844	1	978	MRWACDFTKLLSSTWTLIQYPE
						NAMIIARWALLADCPEKTKAL
						RTTEDGIVDAANFEQFLQERIK
						VNGKVGNLGGGVVTIEKSKSKI
						TVTSKG\PFNSM*I*GSASSFVEK
						KRLLEF*KLLSSTWTLIQYPENA
						MIIARWALLADCPEKTKALRTT
						EDGIVDAANFEQFLQERIKVNG
						KVGNLGGGVVTIEKSKSKITVT
	1					SKGHFQFYVNLRISFLICGKEKA
						VGILEEIVLNLLIIFSSIDILTVKG
ŀ						TCSGCERGFFLEIYDEVEDSSYD
						NHLKDDDPKREGPGAEQALST
						PPCCPLEKRPINDQTLIISWSGST
						NVIEEAGRHSVVQEGWGALGV
1		1				LRTTLPSLPQITGGAGFVGSHLT
		1				DKLMMDGHEVTVVDNFFTGR
		1				KRNVEHWIGHENFELINHDVVE
1		1				PLYIEGRAENQLSHTVPGPVDL
		$oldsymbol{ol}}}}}}}}}}}}}}}}}$				SDRIVNGYFLSL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	- · · · · · · · · · · · · · · · · · · ·
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
18736	49104	A	18845	167	1440	RDIWLSASAAAMVPVRKLV/AE
10750	77107		10045	107		GGQKTKQVLRFTLGCTHPIEDG
						IIDAANFEQFLQERIKVNGKVG
						NLGGGVVTIEKSKSKITVTSKG\
	Ì					PFNSM*I*GSASSFVEKKRLLEF
						*KAENLSSKNWNKTRMPTFTTL
						TQVAPNPITGYTSKGKKISTLKR
						YLAFCLRCCHGAREKACGEGG
	<u> </u>					QKTKQVLRFTLGCTHPIEDGIID
						AANFEQFLQERIKVNGKVGNL
	}					GGGVVTIEKSKSKITVTSKGHF
						QFYVNLRISFLICGKEKAVGILE
			1			EIVLNLLIIFSSIDILTVKGTCSSC
						ERGFFLEICDEVEDSSYDNHLK
						DEDPKREECRKRDAYAPLKDQ
						GNAEKRRHNVRKSWPLLSAKL
						LKIRTNLTGTEEAELHANHIQQ
						VPIKCLSYKLTERKNGTQKHLH
						YALILHVKCDRRETSKFTKFYG
	:	1				THPLTALFVPRAWDGPPSALPE
	}					SELAVFLVAYEGNTLLHLFLSG
	ł	1				AEQRGNSSLKERAGRARFSHQR
						KLNEQTIPPDNLTKSSVDLQSSF
		ł				SAVHPNSQPTVWVDMPAHPLS
						QYLVPARKTVLPIWYGNSCPD
						VIFGSLSDSKLSEGKEFALPLKF
18737	49105	С	18846	713	831	
18738	49106	Α	18847	2	303	
18739	49107	Α	18848	97	233	RTVTCYHRNSRACHPQQITSCP
						*HRALPCLPLWSPKCPSRWPWF
1						G
18740	49108	Α	18849	229	359	MKWVYCSVTGMGKGQHRMP\
						SKG*GWVQRYNRDGIPSPRVLE
		}		j		LF

SEQ ID NO:	SEQ ID NO: of peptide sequence	ı	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
18741	49109	A	18850		1185	MSTIDDPRLQGRRMCLMLRGT PEQKALVIGGEACMWGEYVDN TNLVPRLWLCPPSTQDLANEGI REMTILHRHIPDRGKGVMRSCS FRTPRPFTLACGPEKSLMDPQS QSNSEGASSSLVVKLADTDRER ALRRMQQMAGHLGAFHPAPLP LGACGAYTTAILQHQAALLAA AQGPGLGPVAAVAAQMQHVA AFSLVAAPLLPAAAANSPPGSG PGTLPGLPAPIGVNGFGPLTPQT NGQP\APTRSTITGSPLI/SGWSQ SRPILRIMKYAEQRIPTLNEYCV VCDEQHVFQNGSMLKIQPDTII QVWREDIPVNYMKELELVTKA GFRALLSAPWYLNRISYGPDW KDFYIVEPLSFEGTPEQKALVIG GEACMWGEYVDNTNLVPRLW
18742	49110	A	18851	231	416	AHRIQL
			<u> </u>		<u></u>	
18743	49111	В	18852	184	594	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
					leas (
18744	49112	A	18853	247	2296	PVYLSIFMSCVGENGGSHIENW
						HSILRNGLVNASYTKLQDIKGQ
						FWNDDDSEGDNESEEFLYGVQ
						GSCAADLYRHPQLDADIEAVK
						EIYSENSVSIREYGTIDDVDIAL
						HIN\ISFL\DEEVSTAW\KVLRTE
	1					P\IVLRLRFF\LSQYLDGPDPS\IE
	1					VFQPSNKEGFGLGLQLKKILGM
	1					FTSQQWKHL\SNDFLKTQQEKR
						HSWFK\ASG\TIKKFRAG\LQHSF
						SPIPKS\PSFPYS*Q\DSMAGKGK
						LGV\PELRVGRLMNRSISCTMK
		•				NPKVEVFGYPPSPQAGLLCPQH
	1					VGLPPPARTSPLVSGHCKNIPTL
	1					EYGFLVQIMKYAEQ\RIPTLNEY
	-					CVVCDEQHVFQNGSMLKPAVC
						TRELCVFSFYTLG\VMSGAAEE
						VATGAEVVDLLVAMCRAALES
						PRKSIIFEPYPSVVDPTDPKTLAF
						NP\KKKNYE\RLQKAL\DSVMSI\
			}			REMTQGS\YLEIKKQMDKVWIP
	1					L\AHPLLQWIISSN\RSHIVKLPL
	1	l	}			SRQLKFMHTSHQFLLLSSPPAK
		İ				EARFRTAKKLYGSTFAFHGSHI
		ŀ				E\NW\HSIL/RGNGLVQWHSYTK
						LQLHGAAYGKGIYLSPISSISFG
		l				YSGMGKGQHRMPSKDELVQR
	1					YNRMNTIPQTRSIQS\RFLQSRN
						LNCI\ALC\EVITSKDLQK\HGEH
		ŀ				LGCCPVFRPMSCTRFLPL*YED
		l		Ì		G\QVGDAQHYIPHGPQRVHERE
18745	49113	С	18854	122	486	G (Q V GD/1Q11111 HG1 QRV11ERE
18746	49114	A	18855	1	535	KNAAGNFISPT/ERLSPNAWKTI
10,40	12117	 	10055	•		TDIVLNGTAFVTFLSITTIYAET
	1					GSGFVVPSASAKAGVEAMSKS
	1					LAAEWGKYGMRFNVIOPGPIK
						TKGAFSRLDPTGTFEKEMIGRIP
						CGRLGTVEELANLAAFLCSDY
				1		ASWINGAVIKFDGGEEVLISGEF
10717	140115		10056	2	256	NDLRKVTKEQWDTIEELIRKTK
18747	49115	A	18856	3	356	
18748	49116	Α	18857	1	1581	

SEQ ID S	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
1 - 1	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
se	equence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18749 4	19117	Δ.	18858	2	1242	
10/49 4	+911/	Α	10030	2	1242	1
						CSFCPAPCGPTACGGFRPEFWRI
į						NMKLPAKVFFTLGSRLPCGLAP
						RRFFSYGTKILYQNTEALQSKFF
						SPLQKAMLPPNSFQGKVAFITG
						GGTGLGK\GMTTLLSSLGAQCV
						IASRK\MDVLKATAEQISFSNW
						EIKVHAIQCDVRDPDMVQNTV
:						SELIKVAGHPNIVINNAAGNFIS
						PTERLSP*CWKNQ/ITNIV*NGT
						AFVTLEMGKHLIKAQKGAAFL\
						SITT\IYA\ETGSGFCRYPSASAQ
						SRVWEAHGASLLQLEWG*NM
						GLRIPIVDSNQGPIKTK\GALYPS
				1		GPPTGTFGEKKMIGQKFPCGSP
						WGLVEELRKSLPAF\LCSDYA\S
						W\INGAVIKFD\GGEEVLI\SGDF\
				ľ		NDLRKVT\KEQWDTIEELIRKT
						KRFLRPLWPSSWLQKRE
18750 4	19118	Α	18859	8	432	NSKLPPVVTSQQMRFMY/DPQT
						DQHMKI\FPEQLPLDEFLQKTDP
				ļ		KDPANYILHAVLVHSGDNHGG
				ŀ		HYVVYLNPKGDGKWCKFDDD
						VVSRCTKEEAIEHNYGGHDDD
]		LSVRHCTNAYMLVYIRESKLSE
						VLQAVTDHDIPQQL
18751 4	19119	A	18860	1	211	
18752 4	19120	À	18861	1	242	
18753 4	19121	A	18862	3	733	LVHSRDNHGRHYVVYLNPKGD
						GKWCKFDDDVVS/QLVERL*EE
				,		KRINAQKVDR\RQEVHLYMQA
						QVASEDQFCGHQGNDMYDEE
						KVKYTVFKVLKNSSLAEFVQN
						LSQIMEFPQDQIQLWPMQARSN
						GTKRPAMLDNEADGNKAMIEV
						SDNENSWTIFLETVDLELAASG
				_		VTLPKFDKAHDVMFFLKMYDP
						KMRSLNYCGHIYTPISCEIRDLL
						PVMCNRAGFIQDTSLILYEEVK
					,	LNLTERIQD
18754 4	19122	A	18863	261	659	KHLFYLFIT*EITLYPDKHGCVR
110/37 4	7/122	А	10000	[20]	037	DLLEECKKAVELGEKASGKLR
						LLEIVSYKIIGVHQEDELLECLS
						PATSRTFRIEEIPLDLLEECKKA
						VELGEKASGKLRLLEIVSYKIIG
			1006:		10-	VHQEDELLECLSPATSRTFR
18755 4	19123	Α	18864	107	407	CVFFADTSWRSEATFQFTVERF
						SRLSESVLSPPCFVRNLPWKIM
1						VMPRFYPDRPHQK\SVGFFLQR
1 1						
						NAESDSTEDSLNDATTNPHIVA

SEQ ID	1 ~	1	SEQ ID NO:	1	L	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
10756	140104		110065		12000	VIVIOUGI PROPONI A CARROLA AVA D
18756	49124	A	18865	2	3698	YVRVSLPPPPPAAGRPGAAVAD
						DAREEEEEAAPPPPPPPPRLAA ARPPGSQPRPPAAGEAQAAAD
		1				MNHQQQQQQKAGEQQLSEPE
						DMEMEAGDTDDPPRITQNPVIN
						GNVALSDGHNTAEEDMEDDTS
						WRSEATFQFTVERFSRLSESVLS
						PPCFVRNLPWKIMVMPRFYPDR
						PHQK\SVGFFLQ\CNAESDSTSW
		1				SCHAQAVLKIINYRDDEKSFSR
						RISHLFFHKENDWGFSNFMAW
						SEVTDPEKGFIDDD
18757	49125	A	18866	16	450	
18758	49126	A	18867	3	26	QSGHSMDMRVPAQLLGLLLLR
						LRGARCDIQMTQSPSSLSASVG
			1			DRVTITCRASLDISRWLAWFQL
						KPG\EVPKLLIYAASTLQSGVPS
						RFSGSGSGTDFTLTISSLQSEDIA
		1				MYYCQQYNNW/SSAPYTFGP\G
						TKVEIKTKLWAA\PSVLHLPGPS
			İ			DEAVENLGTAFCLCALLEKTSN
						PQSRPKSTSGRVD\NALPNGGK
						LPQGECSQKQD*TAWT
18759	49127	Α	18868	3	694	
18760	49128	Α	18869	47	1712	FPTKMAAVGRVGSFGSSPPGLS
						STYTGGPLGNEIASGNGGAAAG
						DDEDGQNLWSCILSEVSTRSR\S
						KLPAG*NVLLLGEDGAGKTSLI
						RKIQGIEEYKKGRGLEYLYL\N
		Į.				VHE*MTGNDQTRCN\VWILDG
		ŀ				DLYHKGLLKFSLDAVSLKDTL
		ŀ				VMLVVDMSKPWTALDSLQKW
		ŀ				ASVVREHVDKLKIPPEEMKQM
						EQKLIREF\QEYVERGEDFPASP
						QRRNTASQEDKDDSVVLPLGA
						DTLTHNLGIPVLVVCTKCDAIS
						VLEKEHDYRDEHFDFFQSHIRK
						FCLRYGAALIYTSVKENKNIDL
						VYKYIVQKLYGFPYKIPAVVVE
		1				KDAVFIPAGW\DNDKKIG\ILH\E
		1				NFQTFKRQEDNFWKDIITKPPV
		1				RKFVHEKEIMAEDD\QVFLMKV
						\QSLLSKSKPP/TTEAGRPVDA\S
		1				PKESQGGSPPNTKIESVSS*CCQ
		1				ACHPIPCWGQKKIDPNMESWE
						LTSGRAFLGKFSFNSFV*VKRL
						GSPRRPLVVSGG*PCRVGLGGG
						\SSG\LPP\STKKVQAYKPV\LDV\
L	I		L			HAELDRI\TR*PV\TVS\PTTPTS\P

NO: of pe	سا ددنمت			Nucleotide		Amino acid sequence (X=Unknown,
1			in USSN	location of first		*=Stop codon, /=possible nucleotide
Sequ	ience		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18761 491	29	1	18870	3	885	LVCLVRELLGPRRFFLKVFFLPT
						FPG\VPAPVLTATLPFFHPEGRL
						YQVE\YAFKA\I*PGGPLPSLAV
						RGK\DWAVIV\TQKKST*PNYL
						DSS\TVTSLFPR*LKNIW\CVMT
						GMTS*PADSQV\QRAR\YEA\AN
		- 1				WKIQSMGYEIPVD\MLCKKNCP
						IFLQVYHTEMLKMKASLRC\CM
1						IFNWVIDEEQGPSRYIRC\DPA\G
						YYCGV*KPTASGS*TKLRSTGF
	İ					LEKKKCKKKFD\WTF*TRQWET
	İ					AI\TLPCLTVPIQLDFKTFQEIEV
						l .
						GVVTVENPKF\RILT\EAEIDAQL VAPSKRRD
18762 491	30	4	18871	1	471	VALORIGE
18763 491		-	18872	1	1524	
18764 491	32 <i>A</i>	1	18873	2	719	RPQRAGPVRRAGVMALLDLAL
						EGMAVFGFVLFLVLWLMHFM
						AIIYTRLHLNKKATDKQPYSKL
		1				PGVSLLKPLKGVDPNLINNLET
						FFELDYPKPLFFSVAPMKRIEKT
						QM**P*CSTY*YEVLLCVQDHD
						DPAIDVCKKLLGKYPNVDARLF
						IGGKKVGINPKINNLMPGYEVA
						KYDLIWICDSGIRGTMEMPMSL
		İ				EQSYDPNVRDPIYRNNGDADG
						QHLVLHDFYQQGSGPKCSLINA
18765 491	33 A	4	18874	200	370	
18766 491			18875	323	448	
18767 491	35 A	4	18876	2	421	FVGPGPGDPPPSETHKLVVVGG
						GGVGKSALTIQFIQ/ILDTAGQE
		l				EFGAMREQYMRAGHGFLLVFA
				•		INDRQSFNEVGKLFTQILRVKD
1						RDDFPVVLVGNKADLESQRQV
						PRSEASAFGASHHVAYFEASAK
						LRLNVDEAFE
18768 491		\rightarrow	18877	2	236	
18769 491	.37 A	٩.	18878	1	763	MTPHSALRARGKGGGTGGGSE
						GSSSGGDMSSGAASGTGRGRPR
						GGGPGPGDPPPSETHKLVVVGG
						GGVGKSALTIQFIQSYFVSDYD
						PTIEDSYTKICSVDGIPARLDILD
						PAG\QEEFGAMREQYMRAG/HT
	į					GFLLVFAINDRQEFSTRWGKLF
						HADFCGFKDRD\DFP\VVLVGN
					:	KAD\LESQRQVPRSEASAFGAS\
						HHVAYF\EPSAKMRF\NVDEGF*
						RSWCGVVREIPRNKELPPSPPK
	1					CPPGRMGGGCPCVLL

SEQ ID			SEQ ID NO:		l e	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
18770	49138	A	18879	2	403	RFLARRTNSTFNQVVLKRLFMS
						RTNRPPLSLSRMIRKMKLPGRE
						NKTAVVVGTITDDVRVQEVPK
						LKVCALRVTSRARSRILRAGGK
						ILTF/DEVYRHFGKAPGTPHSHT
						KPYVRSKGRKFERARGRRASR
					ŀ	GYKN
18771	49139	A	18880	1	1006	PKLR*PEPSRNQVLLTRFQIAVH
						F/LLSRLLQFFPRPARSRAFLSRK
						PKPGPGKKPMALIVEFIGELPTG
						VHARPPSHVETLCNTFSSQIEW
						HNLRTDRKGNAKSALALIGTDT
						LAGDNCQLLISGADEQEAHQRL
						SQWLRDEFPHCDAPLAEVKSD
						ELEPLPVSLTNLNPQIIRARTVY
						SGRAGGIQTPISSLDLNALGNLP
						AAKGVDAEQSALENGLTLVLK
						NIEFRLLDSDGATSAILEAHRSL
						AGDTSLREHLLPVGQCANAASS
						TNHPFVTKKRQQCFVFRQRLA
						NGRNDSGWQITHAFWFRFHQH
						SRIMKYLYGFTARLKEHQPKLR
18772	49140	Α	18881	1	716	VCCEFCRAERRHHGEWTIRHN
						K\DRKGSGARDPKEPGFLL*RC
1		1				CVKLYRFSGPEEPNSNIQPRLW
1						LKRLFMS\RT\NRPPSVPFSRDES
				1		GKMKASLARGKQDRPVVVGG
				ļ		PLT*LMLRVSGRVPKTERYVAL
						RRGPSPGPQAAFLQGQGGKIPS
						LFDQAWPLGLPLKGLLALVPGS
						SGSLGKGAREGGTRQFSAKAP
						KETPQKPHQNPYVR\SKGRKFE
10000	10444	 —	10000		1.00	RARGPTGPARGYKKLTLGSYSL
18773	49141	A	18882	3	168	
18774	49142	A	18883	23	2669 396	
18775	49143	A C	18884 18885	960	1250	
18776	49144	+	18886	52	925	EVLEPRLGVFSNGCFQGLSSVM
18///	49143	Α	10000	32	923	ALGLKCFRMVHPTFRNYLAASI
	Ì					RPVSEVTLKTVHERQHGHRQY
						MAYSAVPVRHFATKKAKAKG
						KGQSQTRVNINAALVEDIINLEE
						VNEEMKSVIEALQG*FQ*GLSI*
		1				GPHQDPIDKIAVVTA*REALL*T
						RFSQISMKSPQLNFG*IWASFPE
						CTAAAIKAIRESGMNLNPEV\EG
1						TLIRVPHSPK*PESTEKCWVKL
1						GQTEHQQGPKTLLREGFRTQLN
1		1				EQAGRNPKD/NSLQEDTIRLTEK
						QISQMADDTVAELYRHLAVKT
						KELLG
L	<u></u>	1	<u> </u>		<u> </u>	1100

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		,,,,
18778	49146	A	18887	2	389	DSGSHCHAPPTTARRAFPIPFGS
10770	.,,,,,		1			KSNMATLKDQLIYNLLKEEHTP
		1				QNKITAVRVGADGMACAITIL
]	M/KDYNVTANSKLVIITGGARQ
						QQGESRLNLVQRNVNIFKFIIPN
						VAKYSPNCKLLIRSNPVDI
18779	49147	В	18888	217	423	
18780	49148	Α	18889	27	1217	ARRPTRPPDLIATRPRRPARRCI
						SRFPFGSKSNYWQPLKDQLIYN
						LLKEEQTP\QNKITVVGVGAVG
						MACAISILMKDLADE\LALVDVI
						EDKLKG\EMMDLQHGKPFSFR
						NTKVLSSGKRL*M*LANSKAGS
						FITGWGHRQQEGKKAVLILVQ
						AVNVNIFKFIIP\NVVKYSP\NCK
		1				LLIVSNPVDILTYVAWK\ISGFP
		ł				KNRVIG\SGCNL\DSA\RFR\YLM
		ŀ				G\ERLGSSPH*ACHGWGPWGEH
		1				GRFPVLPVMGVGLNVAGVSLE
ŀ		1				DFWHPDFRTDKD*E\QWKE\VH
						K\QVVE\SAYEVIKLK\GYTSLG
ŀ						LIGLSCKPD\LAE\SIMEESFRRV
						HPS/SSHHGLKGLYGNKRMNVS
		ł				LSVPCILGQNGIS\DLVKG\TLTS
	<u>_</u>	<u> </u>				EGRGPVGSKSADNNSWGSTKK
18781	49149	A	18890	1	264	
18782	49150	Α	18891	3	736	CVCDVA A PIDOCNIEL TODVIEL
18783	49151	Α	18892	81	1065	GKGPVAAFIDQSNIFLTDPKIFL
						GQWREEPKMPLLLLGETEPLKL ERDCRSPVDPWAAASPDLALA
				1		CLCHCQDLSSGAFPNRGVLGG
				1		VLFPTVEMVIKVFVATSSGSIAI
						RKKQQEVVGFLEANKIDFKELD
		ŀ				IAGDEDNRRWMRENVS*REKP
						Q\NGIPLPSQI\FNEE\QYCG\DFD
						SFFSAKKENIIYSF\LGWAPLPDS
		ļ				KGSEKAEEG\GETEAQKEGSED
						VGNLPEAQEKNEEEGETATEET
						\EEIAMEGAEGEAEEEEETAEGE
						EEPGEDEDFLGLFMLHFFHFSQ
						KMEAMKQHSNVLSTNQTETEY
						FGLKPAHPRVTEDIMLPDLIR
18784	49152	A	18893	19	445	TODAL ATTENTION DEIN
10/04	1+7134	l _V	10073	117		

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18785	49153	A	18894	<u> </u>	622	QAAWILKARALTEMVYIDEIDV
10/03	149133	A	10024	1	022	DQEGIAEMMLDENAIAQVPRP
						GTSLKLPGTNQTGGPSQAVRP\
						TSHSEKPITGFLRPSTQSGRPGT
						MEQAIRTPRTAYTARPITSSSGR
						FVRLGTASMLTSPDGPFIN*SRL
						NLTKYSQKPKMAKALPEYIFHH
						ENDVKTALELAALSTEHSQYK
						DWWWKVQIEKRYYRLGMYRE
	l .					AEKQLISAMKQ
10706	40154	A	18895	1	1651	MKAEICKSRTVKDWHNHQKLG
18786	49154	Α	10093	1	1031	EGNKTDSPPQPSEGTNTANTLIL
						DFWPLEMIACCWETFPQRLTAE
						LFMNPTHRWHGYKNQSVGAL
						· ·
						RAPLGQGPSRRGLPGRVGRRQL
						FTPRPPLSWSAGPSLAAPAAMS SEMEPLLLAWSYFRRRKFQLCA
						DLCTQMLEKSPYDQEPDPELPV
						,
						HQAAWILKARALTEMVYIDEID
		1				VDQEGIAEMMLDENAIA/PSSTP
						ITQAGRPITGFLRPSTQSGRPGT
						MEQAIRTPRTAYTARPITSSSGR
						FVRLG\RLGMYREAEKQFKSAL
						KQQEMVDTFLYLAKVYVSLDQ
					1	PVTALNLFKQGLDKFPGEVTLL
		ŀ				CGIARIYEEMNNMSSAAEYYKE
				1		VLKQDNTHVEAIACIGSNHFYS
ĺ					i.	DQPEIALRFYRRLLQMGIYNGQ
						LFNNLGLCCFYAQQYDMTLTS
			• •			FERALSLAENEEEAADVWYNL
		1				GHVAVGIGDTNLAHQCLRLAV
		1				VNSNNHAEAYNNLAVLGDAEG
		1				/RHVEQARALLQTASSISTPYV*
		ĺ				TAFLILQQSLIRFGDLQRSYVAA
						QKSEAAFPDHVDTQHLIKQLRQ
10505	10155	 _	10006		205	HFAML
18787	49155	Α	18896	2	295	MI CLCLI DI DOCCHA VEDCODO
18788	49156	Α	18897	1	4470	MLGLGLLRLRQGSHAVTRCRP
						LPVRREGRRDGSPWRSVVCRY
1		1		1		CRCSRQTGASVTTVSLPSSSSSP
						GLDPRGPRQASVKKPAEANPV
						LRPLVRHGTPYRDSEEGKREGL
1		1				SRLRAVCRRAGPRGRGSFSPRD
		1				ARASPRLHFLVAAVTTGAASRR
1		1				QRGARVRQPSPSSSRRAKRLRE
1				1		CERRSLHAPPAMDASYDGTEV
						TVVMEEIEEAYCYTSPGPPKKK
						KKYKIHGEKTKKPRSAYLLYY
				<u></u>		YDIYLKVQQELPHLPQS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18789	49157	Α	18898	89	810	QRRSGCVLRMTEWETAA\PAG
			[GQRTPDIKALWGSGAPDDVAR
						SMNISLQ\DYIAS*RRKLCPSYLP
						\HSCQGGYAAKNLSRKAQ\CPI\
			į			VER\LTNSPMMMHRPQQPARK
						LMTVRIVQA/HAFEIITPCSQAE
						NPL\QVLGEPPSINSG\PREDSHT
						HLGAAGTVRRQAVDV\SPLRRV
						EPGPFWVACCTRRLVKGLAFPG
						TFKTIG*VAWAD\ELI\NACQGA
						PSNSYCL*EEGRVAFLEHVAKS
						NR
18790	49158	A	18899	1	365	DTHFK/DSHHRTSTAHLL\SLSQ
						GPSPPNLPLGVPIS\GMGLGPRG
	1					EGRANPIPTVLKRDQEGLGYRS
						APQPRVTHFPAWDTRAVAGRE
						RPPRVATLSWREERRREEKDRA
	·					WERDLRTYMNLEF
18791	49159	Α	18900	203	427	
18792	49160	Α	18901	95	482	ALQSPKLWGYFSGRKVFLNYFF
						SSQQVSEAVVATGSPRAWLTC
						LILPLPGIIFSVLPKAMSRPLLITF
						TPATDPSDLWKDGQQQPQPEK
						PESTL\DGLAAKLFYEALIGDES
						SAPDSQRSQTEPARERKR
18793	49161	Α	18902	348	1660	GVTHLFLFGKRKLRNGIAGRIK
						GPGWIFFSYCRSLRLWLLTGSP
						LSLALPCLILPLPGIIFLKFSQKAI
					}	VPGPLLITFTPATDPSDLWK\DG
						QQQPQ\PEKPESTL\DGAAARAF
						YE\ALI\GDESSAPDSQRSQT*T
			İ			WPEERKRKKRGLLKAPAAEAL
						AEG\ASGRPVQGRSLEADDPMT
						YRILTSAQEGD\LP*LM\RLLEPH
						EAGGAGGNINARDAFWWTPL
						MCAARAGQGAAVSYLLGPG\A
						AWVGVCELSGRDAAQ\LAEEA
						GFPEVARMVRESHGETRSPENR
						SPTPSLQYCENCDTHFQD\SNHR
		1				\TSTVHLLSLSQGPQPPQLPLGV
						PISTPRF*ILLLGGVGSPGMGLG
						PRGEGRANPHPQLILK*GPRKG
						LG\YKISTPSPEVTHFPSFGIPRA
						VAGRERPP\RVA\TLNWKE/ERR
						RREEKDRAWERDLRTYMNLEF
18794	49162	С	18903	135	299	
18794	J49162	C	18903	1135	299	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
i	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18795	49163	Α	18904	3	2074	ERSRPVALRAAVPRMNPSTPSY
						PTASLYVGDLHPDVTEAMLYE
		İ				KFSPAGPILSIRICRDLITSGSSN
						YAYVNFQHTKDAEHALDTMNF
		ł				DVIKGKPVRIMWSQRDPSLRKS
						GVGNIFVKNLDKSINNKALYDT
						VSAFGNTLSCKVVCDENGSKG
						YGFVHFETH\EA\AERAIEKMNG
						ML\LNDRKVFVGRFKSRKEREA
						ELGARAKEFT\NVYIKNFGEDM
	ļ					D\DGRLK\DLFGKF\GPALKCEK
		l	ļ			LMTDESGKSKGFGFVS\FERHE
		ŀ				DAQKAVDEMNGKELNGKQIY
						VGRAQKKV\ERQTELKRKFEQ
						MKQDRITRYQGVNL\YVKNLD
						DG\IDDE\RLRKEFSPFGTIQLVP
						KVMDGGLVAAKGLGFVCFSSP
	}					E\EATKAVTEMNF\RIVATEPL\Y
						VALA\QRKEERQAHLTNQYYG
						RGM\ASVTEAVPNPV\IQPLPGQ
						APPFRFTFMGRFSHRFRNRA\AY
						YP\PSQIAQTKTKFLRWTGQGA
		ŀ				RT\HPF\QNMPGCYPAQLAPRPP
		į				I*GTM\RPASSQ\VPR\VMPTQR\
						VANTSNTDKWVPRSCSLQAAA
						AYVL\RPAPFPQYKYAAGVRNS
						\QQHL\NAQPQVTMQQPCCSMV
						PRVQGTLWTCLPMVGHLPPPSR
						SKSQMLGWNGLFPSYSKPMHP
						YSLLVKSLG\MLLEIDNSEL\LH
						ML\ESP\ESLPF*RVDEAVAGTTS
				}		PPQAKRGLAQKAG*QVPTGVP
18796	49164	Α	18905	1	988	MAEDGEEAEFHFAALYISGQW
						PRLRADTDLQRLGSSAMAPSRK
		ļ				FFVGGN\WKMNGR\KRRLGELI
						GTLNAAKVPADTEVVCAPPTA
						YIDF\ARQK\LDPKIAVAAQNCY
						KI\TN\GAFTGEISPG\MIKDCGQ
						PWVVLGHSERRHVF\GESDELI\
						GQKVA\HALAEGL\GVIRLHLG
				1		RS*DERGSLGIT\EKVVFE\QTKV
				1		IRR*T*KDWKQ/VVVLGLMSPV
						WGLWLLARLATPQQA\QEVHR
						RSSRGWL\KSNVS\DAVASEPPV
				1		TIYG\GSV\TGATLQRSLAKPSP
				1		DVDGLPLWGGASLQARNSWDI
						IQCHNNEPPSHLSLPLRCQAQG
I	Į.		l			LKQTQKPK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
18797	49165	A	18906	46	468	TGPTHASAAILVPRSLHKMPGE
						ATETVPATEQELPQPQAETGSG
						TESDS/GKSPASDTYIVFGEAKIE
						DLSQQAQLAAAEKFKVQGEAV
						SNIQENTQTPTVQEESEEEEVDE
						TGVEVKDIELVMSQANVSRAK
						AVRALKNNS
18798	49166	Α	18907	1	360	
18799	49167	Α	18908	52	513	QLGPHGHRQGPGGPAVPHPRG
						LVPLPPALLPSGPRRGRTQAPR
						HSGPAL*GHPASDTYIVFGEAKI
						EDLSQQAQLAAAEKFKVQGEA
						VSNIQD\NTQTPTVQEESEEEEV
						DETGVEVKDIELVMSQANVSR
						AKAVRALKNNSNDIVNAIMEL
18800	49168	Α	18909	1	422	RSLCQTGLRQ/VTG/VTKVTIGN
						LENLFVI/TKPDVYKSPASDTYI
						VFGEAKIEDLSQQAQLAAAEKF
						KVQGEAVSNIQENTQTPTVQEE
						SEEEEVDETGVEVKDIELVMSQ
						ANVSRAKAVRALKNNSNDIVN
10001	10160	_	10010	107	500	AIMVSVQAFVP
18801	49169	В	18910	427	529 252	
18802	49170	A	18911 18912	122	189	SKLDYKSSKRHRKPVSFNFITSF
18803	49171	Α	10912	44	109	QIWD*FLFFF*LKCQLQPRVFSQ
						PN
18804	49172	Α	18913	2	609	PFLPEAHLGNQKCQNCNSTSGI
						CRVSVKVCPGPWVHAKLCAPT
						LMSRDVTLPDAPVKKPCGEGG
						AKKKKQVLKFTL\DCTHP\VED
						G\IMDAAQF*ASFCKERIKV\NG
						KAGNL\GGGVVDPSKGARGKIT
						VT\SEVPFSKRVF*KYL/TPKKY
		1				LKKNNLR*PGLPRSLPNQPKEK
						LPNFRYFQINQGRRKEGGRRGF
10005	40150	ļ	10014	100	1504	KFPLSGKILL
18805	49173	A	18914	107	524	AGTHLPPFLPVAVRQPMALR\Y
						P\IAVGLKQGPTKVTQEP*ASPR
						HHVRRGRL\TKHTKFVRDMIRE
						VCGFSP\YE\RRAMELLKVSKD
						K\RALKFIKK\RVGTHIRAKRKR
						EELEQQYWPAIEEKLACQERLE
10006	10171		10015	20	165	PLPLPSPLK
18806	49174	A_	18915	39	465	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
18807	49175		18916	439	1586	TWGPPARRLPSGSHVPSA/ASSR LPSHPREEPQLPQPAPVTITATM SSEAETQQPPAAPP\GAPALSAA DTKPGTTGSGAGSGGPGGLTSA APAGG\DKKVIATKVLGT\VKW FNVRNGYGFINRNDT\KEDVFV HQTAIKKNNPQKYLRSVGDGE TVEF\DVVEGEKGAE\AANVTG\ PGGVPVQG\SKYAADRNHYRR YPRRRGPPRHYPQ\NYQNS\ESG EKNRGIGRVLPEGQAQQRRPY\ RRRFPPYYMRRPYG\RRPQY\S NPPVQGEVMEGADNQGA\GEQ GRPVRQNMYRGYRPF\RRGP\ PRQKTA*ERAAMEEDK\ENQGV WDPRFGSPPQ\RRYRRNF\NYR\ RRRPQKTPKP\QDGQKRPKAAG SHPA*EFVPLPEAE\QGGAE
18808	49176	Α	18917	1	651	MTAFNSGKVDIVAINDPFIDLN YMVYMFLYDSTHGKFHGTVK AENGTLVINGNPIT\IFQDQDPS KIKWAP/LAKVIHDNFGIIEGFM TTVHTITATQTINGPSG\NCHVM AAGLSRTSSLSLLA/LAKPVGK VIPELNGKLTGMAFHVPTANVS VADLTCRLEKPA/KYDDIKK\NT HSSTFDAGAAIVLKDHSVKLIS WYDNEFGYSNRVVHLMAHNA
18809	49177	Α	18918	2	5264	

SEQ ID		ł .	SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
			03,210,21	sequence		, , , , , , , , , , , , , , , , , , , ,
18810	49178	Α	18919	3	1466	PRSAPRAASWLEDPREVRSVCL
						SAPPVRQSAAIFFCVASRATSLR
						TPMGKVKVGVNGFGRIGHLVT
						RAAFNSGKVDIVAINDPFIDLN
						YMVYMFQYDSTHGKFHGT\VK
						AENGKLVINGNPITIFQE\RDPSK
						IKWG\DAGAEY\VVE\STGVFNQ
		1				PWRKAGAHLQGG\AKRVI\ISAP
						SG*MPPMFVMG\VNHEEV*QTA
			:			FKIISN\ASCTNQLA*QPLAKG*F
						HDNFG*SWEGLMTTVPWPSLG
				-		NPRKTVD\GPSRG\NCGPWMGR
			i	-		GGFSRNIFLCLYWALPRAVGKG
						HP*A*TGKLTG\MAFPVSPTAN
						VVSGWTLTCRFRKNLPKY\DDI
						KEGW*KQAFGGAPSKGIPGLQL
		ŀ				SHQVV\SSDFNSDT\HSFHPFDA
						GA\GIA\LNDHFVKL\ISWYDNE
						FGYSNRVVDLIGPHGLQGSKTP
ĺ						WTTEPQARSKKGKERPSLLGSP
						CHTQTPTTLNLPSSQLPCRPLEE
ŀ						GRGLWEPHLVMYPSNKVSLCS
						TKKKKKKKKP
18811	49179	Α	18920	3	364	STVINIHSETSVPDHV/VLVPVQ
						HPLLE\RCCLGFIAFAYSVKSRD
		İ				RKMVGDVTGAQ/ALCLHRQVP
						EHLGPDSGHPHDHWIHPVTGIR
		İ				LCDSLPYYVTDNTGKTGLLVA
						AHSLQPLHSTVQCW
18812	49180	Α	18921	41	696	PDRRWSSLGHHESHCPNHSSLL
]						STVASPPNYEDAQGR*SR/EVA
						VLGAPHNP\APPTSTV\IHIRQSR
						PPWPDHVVWSRVQHPLQ*TPA
						C\LGFIAFAYSVKSRGQGRWLG
						DV\TGAQGLMPPPAKVPETIWG
				<u> </u>		PDFWGILNDHF\VHRHPQWLIFP
						G\HGIDQGRHHWRPGALPI/VTC
			1			IPRLLPT\SIPRPAPRSPKSCISPYI
						LTRFSTMAFNKSARVSGKKKK

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
18813	49181	A	18922	127		SNLILRNLFCPHLALASSAFDKC QEEETVEKTELLAERRGLTENH PGNPITAGHHEPHCANLLSCQQ RPRLPNYEMLKEEQEVAMLGA PQKPCLPECPPVIHYPQARPPWP DHVRLGPCFNNPLQ*TTCCPGL SLGISAYFRGSFRGTGKMV\GD VEPGPQGLMPPTAQVAP*TSGA PDFLGSLHEPILAHQSSPPKCW VVPGPAIDQE\ASLRP\GA\LPVT C\IPRTL\SSIPSPLPPEARELLPFD LYSTYSTFHSSPCPHSRVLHQPF ILTRFSTMAFNKVYMFLVKKK KKKKKKV
18814	49182	A	18923	115	652	VSESTDLVSIGRLVTRA\AFNSG KVDIVAINDPFIDLNYMVYMF\ QYDSTHGKFHGTVKAENGKLV INGNPITIFQE\RDPSKIKWGDA G\AE\YVVES/TLGVFTTMEK\A GGSFCRGGAKRVIHLLPPLA*C PPCFVMGVN\HEKDDN\TLKIIS N\ASCTTN\CLAPLASTGAAKGC GQGHPEA
18815	49183	A	18924	91	399	
18816	49184	Α	18925	26	304	
18817	49185	Α	18926	18	203	
18818	49186	Α	18927	23	355	
18819	49187	A	18928	I	211	RLLRVRGPPSPRPPGTPAGRQA RHTAPVPARASPSTPPRKLKER APALASPERGSHSAAEG*RAPQ VPPK
18820	49188	Α	18929	2	800	
18821	49189	A	18930	1098	1560	VSCPFTVIIFCYNFCRGGFSTTA VMARAKSMFLMLKGKAGPDF GIRFTASSGWFT*FEDRYALHN VNMNAESVGADVKAAE\DFGS LIMEEIYLPTQFFNMEEPSLFWK QMPERTFVICPKGFFPDHCNKE RPWHFCKRKSLTDTRVATPCG
18822	49190	A	18931	33	211	
18823	49191	А	18932	110	624	CILSKMLRSPQELARLPGGQRQ LTRW*QRWPLWTIALVLVLWR VKSGP*S*RVPSPSP/PVEPWVT LNMAGAPHGNPAFPPGGPPHP VPQPGYPGCQPLGPYPPPYPPPA PGIPPVNPLAPGMVGPAVIVDK KMQKKMKKAHKKMHKHQKH HKYHKHDKHSSSSSSSSSSSDSD

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18824	49192	IA	18933	2	664	CEPLRISGRPTRPGERSKPRRRL
						ER*WGWRTHLNMWNPNAGQP
						GPNPYPPNIGCPGGSNPAHPPPI
						NPPFPPGPCPPPPGA\PHGNPAFP
						T\GGPPHPVPQP\GYPG\CQPLGP
		1				YPPPYPPPAPG\IPPVNPLAPG\M
						VGTKQLISRQEDCRRKLKKAS*
						KRCHKA/HQKHHK\YHKHGKH
						FLLFPPPSFQAVISGLEYRALDP
		İ		-		SLKSHQFCSPIKLQMPCCTGGM
18825	49193	Α	18934	3	409	QTQVVIDAG/ALAVFPSLLTNPK
						TNIQKEATWTMSNITASR/QDQI
						QQVVNHGLVPFLVSDLS/KDFE
					:	TQKEAVWA/GGTVEQIVYLVH
						CGIIEPLMNLLPAKDTKIILVILD
						AISNIFQAAEKLGETEKLSIMIEE
						cggv
18826	49194	Α	18935	133	400	VLGPHGLGDHLGQFHSCQSRA
						EPPIAGEDIHAGLDEPDCLTKDF
					-	LRI*LQLSTQLHAGQEEEDQNV
						VPETTSEGYTFQVQDGAPGTFN
18827	49195	Α	18936	218	863	
18828	49196	Α	18937	106	1879	TQSALQPLIPPASPSPWKAQARF
	İ	ŀ				GAFSLCLITMSTNENANTPACP
						VVHRFKNKGKDSTEMR\RRRIE
						VNVEL\RKAKKDDQMLKRERL
						*AHFLDDATSPLQEN\RNNQG/T
						LLNWSVDDIVK\GINSS\NVENQ
						LQATQAAR\KLLSREKQPPIDNII
						RAGLIPKFVSFLGRT\DCSP\IQF
		ŀ	ł			ESAWALTNIASG\TSGQTKAVV
						DGGAIPAFIFSGWASSHAHISEQ
	ľ	1				AVWALGNIAGDGSVFRDLVIK
				1		YGAVDP\LLALLASS**CQSLAC
						GYLR\NLT\WTLSNLC\RNKNPA
				ĺ		PP\IDAVEQILPT\LVRLLHHDDP
						GSVRQDT\CWGYFPTLLDGSK
						WNGIGHGWV/VKQGVVPQLVK
		Ì				LLGASE\LPIVTPA\LKSP*GNNW
						HLGTDEQTQVVD/IDAGALAV\
						FPSLLTNPKTNIQK\EATWTMSN
						ITAGRQ\DQIQQV/V*NHGLVPF
						PLSVFLSKADFKTQKEA\VWAV
						TNYTSGGTS*TELVYLV\HCG\II
						EPLMNLFYCKRLPRIILLILDAN
						FKISFQPAENLV\DLEKPSINLE\
						ECGGLRPKLKALPKPMKMESV
						YKASVKA*FEKYFPCRGKEGRI
L						KTVVP\ETT\SEGYTFPSSRMGPP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
18829	49197	A	18938	12	251	KVTNVKDGTTH\QTSLELFMYL
						NEVAGKHGVGRIDIVENRFIGM
						KSR/GMNVQGDYEPTDATGFIN
						INSLRLKEYHRFQSKVTVK
18830	49198	Α	18939	1	888	
18831	49199	Α	18940	1	1155	MSSKGSMVLAYSGGLNTSCILV
						WLKEQGYDVTAYLANTGQKE
						DFEEARKKALKLGAKKVFIEDV
		1				SREFVEEFIWAAIQSSALYEDH
						YLLGTSLTRPCIAHKKVETVQR
						VGAKYEGPWRAPWRMPEFYN
						WFKGRSDLMEYAKQHGIPILVT
						PKNLWSIDKNLMHISYKAGILE
						NPKNQVPPGLYTKIQDPAKAPN
						TPDILKIQFKKGVPVRVTSVKD
ĺ						GTTHQTSLELFMYLNEVAGKH
						GVG/YIAIVENIFTRQRP\SPECEF
						VRPCIAKSQEPAEGKVQVPVLK
						GQVYILGWESPLSLYNEELMSV
		ŀ				NVQGDYEPIDDTGFININSLRLK
						EYHCLQSLLEFAGCRLQTLFAW
						VSPAGAAEQQRLLPVPSSGNFIP
						EGHLPDASWSSPV
18832	49200	Α	18941	1	993	MSSKGSVVLAYSGGLDTSCILV
						WLKEQGYDVIAYLGQAVKGN
						QLLPVSLVKRKTTPGAQYANR
						LSPRVGRFINAAGTTGFPTGKR
						AGNDQVRFELSCYSLAPQIKEM
						GSLKVRNE/LQEVSTVPMPAST
	ı					QLPADTQCKLAE/LSAPVKK*T
						LGLSKT*QTQVMSSQSADSMPR
						AWYRN*LFTNPTVRRLGNSPVT
						GSSLPRLPG/RGIYETPAGTILYH
						AHLDIEAFTMDREVRKIKQGLG
						LKFAELVYTGAECKLLVASVPE
						PWRGRGGVEEGGRDIKEVQEH
						GSDSAESSTHPPASRPGEGCLA
						CIFCGKWRHLNTPLSCERYPRH
						SRDHRPH
18833	49201	В	18942	324	1035	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18834	49202	Α	18943	29	1516	KQIPDAGNSRLQSQTLLSSKGS
						VVLAY\SGGLDTSCILVWLKEQ
						GYDVIAYLANIGQKEDFEEARK
						KALKLGAKKVFIEDVSREFVE\E
						FIW\PAIQVQAHLY\EDRYLPGH
						LSLPGPCI\ARKQVGNSPSGRGA
						KYVVPTGATGKG\NEQV\RFEL
						S\CYSL\APQIKVIAP\WR\MPEFY
						NR\FKGRNDL\MEYGKATLGFPI
						PVTPKNP\WSMDENLMHI\SYE
				1		AGILENPKNQAPPGLYTK\TQGP
						SQKPPTPLDILGDSSFKKGVPVE
	ļ			1		GGPTFKDG\TTHQTFL\ELF\MY
						LNEVAG\KHGRRAVFDIV\ENRF
						\IGN*SPRGILRRTP\AGHHSFYH
						AHLDIRGLSPWDRGSCGKIKQG
						PGAWKFAEL\VYTGFPAQAPE\
ĺ						CEFCPPTGIAKVPRKPSGKGKV
						QVVRSFKGPQVYI/LSGREIPHC
				i		LLLTMRELVKHGTCKGDYWRP
						IDGHRGSSKHQIPFKAEGNYHR
į					İ	LPRAKVTCQIDPVYNEELGPPQ
						FSGSPQVTGANCCDNL
18835	49203	Α	18944	1	560	GRARTPANMALRVVRSVRALL
						CSLRALWAPAVPCLPRPWQLG
						AGAIWTLLTGPVLLWVCKFTE
						KHEWITTENGIGTVGINNFANE
1						ALGDVVYCSLSEVGTKI*TKPS
						WSLVLLE\SVKAASEL\FSPLSG
					1	EVPDFNEALAE\NPG\LVTKSRY
						EDG\WLIKM\TLSNPFRT*MHL
						MSEEAYEKYIK\SIEE
18836	49204	Α	18945	3	108	
18837	49205	Α	18946	1	845	MAIPGIPYERRLLIMADPRDKA
						LQDYRKKLLEHKEIDGRLKELR
						EQLKELTKQYEKSENDLKALQS
						VGQIVGEVLKQLTEEKFIVKAT
İ						NG\PRYVVGCRRQVIELPLTNPE
						LFQRVGIIPPKGCLLYGPPGTGK
						TLLARAVASQLDCNFLKVVSSS
						IVDKYIGESARLIREMFNYARD
						HQPCIIFMDEIDAIDIDLPNEQA
						RLDILKIHAGPITKHGEIDYEAI
						VKLSDGFNGADLRNVCTEAGM
						FAIRADHDFVVQEDFMKAVRK
L		<u> </u>				VADSKKLESKLDYKPV

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
18838	49206	A	18947	3	1181	LLIMADPRDKALQDYRKKLLE HKEIDGRIKELREQLKELTKQY EKSENDLKALQSV\GQI\VGEVL KQLT\EEKFIVKATNGPRYVVG C\RRQL*QKVSLKPG\TRVAL\D MTTLTI\MEDILPR\EVDPLVYN MS\HEDPG\DVSYSEIGGLS\EQI RGI*GEVIGLPLYKPQSYFQRVG IIPPK\GCLLYG\PPGTGKNT/LLA RAVASQLDCNFLKVVSSSIVDK YIGESARLIREMFNYARDHQPCI IFM\DEIDAIGG\RRFSEGTSADR EIQRTLMELLNQMDGFDTLHR VKMIMATNRPDTLGP\ALLRPG RLDRKIHIDLPNEQARLDIL\KIH AGPITK\HGEIDFESNC*SFRMG FNGADL\RNVCT*SQACSPIRAD
18839	49207	A	18948	1	220	P*FC*YREDF\MKA\VRKVA VNMRDRFGQIMIENLRRRQCEL AGVETCKSLES\RIESLEFLDEM */LLEQLM/RHYCLCWATKGGN ELGLKEITY
18840	49208	Α	18949	1	645	
18841	49210	A	18951	10	502	AAPAEPGRALPSSVAFSLWLAP SPAARRPRFHVPGGAQLPGTVH ARWPARQR\ESSI\TSCCSTSSCD ADDEGVRGTCEDASLCKRFAV SIG\YW\HDPYIQDL\VRLSKQR\ KAPEINRGYFARVHGCSVQLIK\ AFLR\KTECHCSNCSNLGGQGM GSPPFW\RLKDEDLSSQVNIFEV DFPMIVT\RKLHSIKWLAFPLSS PILELHSEDTLQMASDCIC\DGH ILDSKRYAVIGADLRDLSELEE KLKKCNMNTQL\PTP*IAECVL VYMTPE\QSANLLKWAA\NSFE RAMFI\YYQQVNMGDLFGQIMI ENLRGRPV/CDLAGVET\CKSLE SQKERLLS\NGWENKHRPVRT* LEFVPPGLPSKLK*SRIESLEI\LD ENWELLEQLIP\HYCL\CWATQR RK*SLGLKEITY
18842	49210	A	18951	3	502	TEANTLN*RGLQNME/ARLAE/ RKFM/NPFNMPNLYQKLESD/P RTRTLLSDPTYRELIEQLRNKPS DLGTKLQDPRIMTTLSVLLGVD LGSMDEEEEIATPPPPLPSSSMS LSPAPSCCLGCSPIVGFFFIWGS GHVMGRGGGSSSLRSQLSHVV YSASPSPIKQASWAWL
18843	49211	В	18952	1	1071	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18844	49212	В	18953	67	1026	
18845	49213	A	18954	2	1910	PVASSMFQKAARAVGNAERTD
						SIQRCSGPRCAMEQVNELKEKG
						IKALSVGNIDDALQ\CYSEAI*/R
						LDPHNHVLYSNR\SAAYAKKG
			 			DYQKAYEDGCKTVDLK\P\DW
						GKGYSRKAAALEFLYRFEEAK
		l				RTYEEGLKHEANNP\QLKEGLQ
						N\MEARLAK\RKFMNPFNMPNL
		1				YQKLESDPRTRTLLSDPTYRELI
		l				EQLRNKPSDLGTKLQDPRIMTT
						LSDLLGVDLGSMDEEEEIATPA
		l				PPPPAKKETKPEPMEEDLPENK
						KQALKEKELGNDAYKKKDFDT
						ALKHYDKAKELDPTNMTYITN
						QAAV\WFEKGDYNKCRELCEK
						AIEVGRES\REDYRQIAKAY\ARI
		1				GNSYFKEEKYKDS\IHF\YNKSL
						AEHRTPDVLKKCQQAEKILKEQ
						ERLAYINPDLALEEKNKGNECF\
		1				QKGDYPQ\AMKHYTEAI\KRNP
						KDAKL\YSIRAACYTKLLEFQL
						ALKDWEEWYQLEPTFIKGYTR
		١.				KA\AALEA\MKDYTKAMDVYQ
		l				KALDLDFSCKEAADGYQRCIM
						AQYNRHDSPEDVKRR\AMADP
						EVQQFMS\DPA\MRFILKQMQK
						DP\QALSE/HL*RNPLLPREIQKL
]		MDVGSDCKFGDDLFIPAFLCPS
						CGKRSWDRGEQAARSGRESST
100:5	1001:		10055		1000	ERKGEQGERRPSSPYIYT
18846	49214	A	18955	2	293	
18847	49215	В	18956	53	182	
18848	49216	A	18957	2	403	
18849	49217	A	18958	3	1646	
18850	49218	A	18959	3	408	
18851	49219	Α	18960	1	2458	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	•	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
18852	49220	Λ	18961	88	1511	LWPLGMASNSSSCPTPGGGHL NGYPVPPYAFFFPPMLGGLSPP GALTTLQHQLPVSGYSTPSPATI ETQSSSSEEIVPSPPSPPPLPRIYK PCFVCQDKSSGYHYGVSACEG CKGFFRRSIQKNMVYTCHRDK NCIINKVTRNRCQYCRLQKCFE VGMSKESVRNDRNKKKKEVPK PECSESYTLTPEVGELIEKVRKA HQ\ETFPALCQLGKYTT\NNS\SE QRV\SLDIDLWDKFSELSTKCII KNGEGRQGRLPGFTTLTI\ADQI TLLK\AACL\DILILRICTRYTPE AGTP*PSWDGLTLNR\TQMHNA G\FGPL\TDLVFAF\ANQLLPL\E MDD/ARETGLLSAICLICGDRQ DLEQPDRVDMLQEPLLEALKV YVRKRRPSRPHMFPKMLMKIT DLRSISAKGAERVITLKMEIPGS MPPLIQEMLENSEGLDTLSGQP GGGGRDGGGLAPPPG\SCSPSLS PSSNRSSPATHSP

SEQ ID	_		SEQ ID NO:	1		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				Sequence		
18853	49221	Α	18962	143	2159	LTHSSTWLGRSCNHDRRQMGS
						KVMSYVVAGRGTCAQETPIYT
						TIRYLEDKEQYGGSLGVDSVLA
						RLFSGASVLFISGPPFSSYLSGVP
					<u> </u>	PGHPMEDSM\DM\DMSPL\RPQ
						NYLFGCELK\ADKDYHFKVD\N
ì						DENE\HQLSLR\TVNLGGLVAK
						G*VAHLLKAEAIEFTEASPNLK
		l				VTTWAT\LKM\SVPAQRVFPLG
						GLLKITPTRWVLKGLKVWVQG
						PVHISGTAL**LWEGRCQSPEDE
		l				E\EEDVNRVLFMKPKGRGLKH
						MFGDLVCSWKLAAIETQSSSSE
						EIVPSPPSPPPLPRIYKPCFVCQD
						KSSGYHYGVSACEGCKGFFRRS
						IQKNMVYTCHRDKNCIINKVTR
						NRCQYCRLQKCFEVGMSKESV
		ļ				RNDRNKKKKEVPKPECSESYTL
						TPEVGELIEKVRKAHQETFPAL
						CQLGKYTTNNSSEQRVSLDIDL
						WDKFSELSTKCIIKTVEFAKQLP
		l				GFTTLTIADQITLLKAACLDILIL
						RICTRYTPEQDTMTFSDGLTLN
		ŀ				RTQMHNAGFGPLTDLVFAFAN
		1				QLLPLEMDDAETGLLSAICLICG
		l				DRQDLEQPDRVDMLQEPLLEA
						LKVYVRKRRPSRPHMFPKMLM
						KITDLRSISAKGAERVITLKMEI
		l				PGSMPPLIQEMLENSEGLDTLS
		l				GQPGGGGRDGGGLAPPPGSCSP
		i				SLSPSSNRSSPATHSP
18854	49222	Α	18963	748	1075	ILPTSLFFLFCFVFFVCF*DRVLL
						LSPG\WSAVARSWLYCNLSLRG
						FKGFSCLSLLSNWDYRCTPLRS
						ANFVFL/CRDRVSPCWPTSVSNS
						*PQ\VIHPPWPPKVLGITRV
18855	49223	A	18964	3	674	GRRRLLRDAEGPEETVRLWPA
	.,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,					ARAAMDAAEVEFLAEKELVTII
						PNFSLDKIYLIGGDLGPFNPG\LP
						VEVPLWL/ARLNLKQRQK\CRL
						LPPEWMDVEKLEKMRDHE\RK
						EETF\TPMPKPFTTWELT\KLL\L
1						NHA\SDNIPKGRTEFRDPGSRVF
1						WGHSY*AKFRVFG\DSF\VRQQ
						EAHAKLD*LGPWMGDQQPAVT
						FLTQA\LNPHVTNSRTNPPSPLE
1						STSVLRTS
Щ.	L	L			l	DISVERIS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18856	49224	A	18965	261	876	VAQKIDMPEYQGEPDEISIQKC
10000	17227	A	10703	201	0,0	QEAVRQVQGPVLVEDTCLSFN
						ALGGLPGPYIKV/WFLEKLKP\E
		1				GLHQLLA\GF\EDKSA\YALCTF
						CTQAPGDPS\QPV\RLVSGGRDL
		1				
						RGRIRWHPEGCQ\DFGWD\PC\F
						QPDG\YEQTYAEMP*GGRRNA
		1				V\SH\RFRAPCWEPAREYFWQV
						WPSLTSASWKEGPSRPGDLGK
		 	10011		1.01	GLAPKPSPHRGRAPP
18857	49225	Α	18966	2	181	TAEKMAVLAPLIALVYSVPRLS
					;	RWLAQPYYLLSALLSAAFLLVR
						KFPPLCHGLPTQREDGNPCDFD
						WREVEILMFLSAILMMKNRRSS
		<u> </u>				TK*AHPTRRR
18858	49226	Α	18967	5331	5454	ETEA*RINDLL*LLEFFCNI*QN*
						P*IRPTVIRSSLKYNT
18859	49227	Α	18968	5186	5308	ETEA*RINDLL*LLEFFCNI*QN*
						P*IRPTVIRSSLKYNT
18860	49228	Α	18969	4570	4691	ETEA*RINDLL*LLEFFCNI*QN*
						P*IRPTVIRSSLKYNT
18861	49229	Α	18970	5666	5788	ETEA*RINDLL*LLEFFCNI*QN*
						P*IRPTVIRSSLKYNT
18862	49230	Α	18971	5630	5752	ETEA*RINDLL*LLEFFCNI*QN*
						P*IRPTVIRSSLKYNT
18863	49231	Α	18972	224	456	
18864	49232	Α	18973	5543	5665	ETEA*RINDLL*LLEFFCNI*QN*
		Ì				P*IRPTVIRSSLKYNT
18865	49233	Α	18974	4602	4725	ETEA*RINDLL*LLEFFCNI*QN*
		}				P*IRPTVIRSSLKYNT
18866	49234	Α	18975	3	313	
18867	49235	Α	18976	5480	5602	ETEA*RINDLL*LLEFFCNI*QN*
		ŀ				P*IRPTVIRSSLKYNT
18868	49236	A	18977	269	394	
18869	49237	Α	18978	5322	5445	ETEA*RINDLL*LLEFFCNI*QN*
		ŀ				P*IRPTVIRSSLKYNT
18870	49238	Α	18979	164	434	
18871	49239	Α	18980	5364	5487	ETEA*RINDLL*LLEFFCNI*QN*
						P*IRPTVIRSSLKYNT
18872	49240	В	18981	18	1904	
18873	49241	A	18982	1296	1973	RILPGGQQNSSWNFSEDTVISIL
						NTINEVIAENLEAAKKLRETQGI
	}	1				EKLVLINKSGNRSEKEVRAAAL
						VLQTIWGYKELRKPLEKEGWK
						KSDFQVNLNNASRSQSSHSYDD
						STLPLI*PGPKNQIRNLIGKKFR*
		1				AIWDQTQNH*IT/DYSTPNERGD
		1				HNRTLDRSGDLGDMEPLKGTT
		1			1	PLMQDEGQESLEEELDVLVLD
						DEGGQVSYPSMVCPSVTPKIVL
L	L	<u>L</u>	<u></u>	1	1	EEGGS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
18874	49242	A	18983	5844	5967	L ETEA*RINDLL*LLEFFCNI*QN*
	1	`				P*IRPTVIRSSLKYNT
18875	49243	A	18984	5781	5904	ETEA*RINDLL*LLEFFCNI*QN*
10075	1,32,13	-	1050.			P*IRPTVIRSSLKYNT
18876	49244	В	18985	1	1477	
18877	49245	Α	18986	152	1539	
18878	49246	Α	18987	5571	5694	ETEA*RINDLL*LLEFFCNI*QN*
		<u> </u>				P*IRPTVIRSSLKYNT
18879	49247	A	18988	4551	4674	ETEA*RINDLL*LLEFFCNI*QN*
	100.10	<u> </u>	10000			P*IRPTVIRSSLKYNT
18880	49248	Α	18989	2	1043	GVRLRYSPIAVVMVGEAGRDL
		1				RRRRAVAVTAEKMAVLAPLI/S
						LSVYSVPR\LSRWLAQPYYLLS
						A\LLSAAF\LLVRKLPPLCHGLP
		1				TQ\REDGNP\CDFDWRE\VEILM
						FLSAIVMMKN\RRSITVEQHIGN
ŀ						IFMFK/SKVANTILFF\RLDIRMG
ľ	1]	LLYITLCIVFLMTCKPP/LSPYM
]	GPEYIKYFN\DKPFDEELERDNR
						VNW/ILLEFFGNWSNDWQSFAP
						IYADLSLKYNCTGLNFGKVDV
						GRYTDVSTRYKVSTSPLTK\QLP
	1					TL\ILFQGGQGGQCGRPQI*QGK
		1				GR/ALFSWDLL*GT*KEGQENVI
	}					REFNLNELYQRAKKLSKAGDNI
		<u> </u>	10000			P\EEQPVASTPTTVSDG\ENKKD
18881	49249	A	18990	1	1105	MESTGEDAVNIVEMVTEDLVY
		ŀ				RLVVLKSLQVWLSPGDFMGSE
		1			•	GTECMLIGPWVAMGRPSESTLS
		1				SHSGPRSPHGTGSPAPRLQAVH
						VLKCPKSRGVQGGWGLASQCL
						PEHRHTQLSCDSTQAWPQLCSA
						QSLRHNRSGQGAGAVTSEPAG
ŀ		1				AGGLPRPPRTQGCLGAES/PAG
		1				QLHSQQRGGPEEGRLLSDVDHP
						DLSSSQQRGPEGP/QVAGGWCV
		l				SAAWSSTHLAGPQQRWDQEEA
						RQWEQIPLSLLMMGDPPGYSR
						APGDPGPQPQPRQLQLCLGSSH
						PTNS*GAELPPVPGSAECAALT
						VLPSLHLPQWWQVGRPGPT*V
		1			1	DPRVGLPGQQAPEQRVYSLTHP
10000	10050		10001	105	202	EQLPVLQVPLLIYRYSSTYNGV
18882	49250	C	18991	105	393	I VWOLDKCKI I DBII TOCIVAD
18883	49251	Α	18992	38	230	LYWQKRKGKLLNNLTQGIVAD
						PVR*KKFHFKCPSNCPKIHNTES
10004	40252	-	19002	197	747	AYAVSFMKYSAHVLGSLYEM
18884 18885	49252 49253	A C	18993 18994	179	747 649	
18885	49253	A	18994	3	3187	
18887	49254	A	18995	274	423	
1000/	77433	17	10790	1217	747	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
18888	49256	A	18997	1333	2243	RCWRPRTLGFASPPTRPCSPTH APGCEMTPRDWHRVLSFASPLR IEEIFQGYDKTFGLKNKKGAKQ QKFIKAVTHQVKFGQ/SKSTSGS TE*SWKRKLKKDDKKKELQEL NELFKPVVAAQKI/RVKGADPK SVV\CAFFKQGQCTKGDKCKFS HDLTLGEKM\EKRSVYIDARDE ELEKETKSSLIQADTMDNWDE KKLEEVVNKKHGEAEKKKPKT QIVCKHFLEAIENNKYGWFWV CPGGGDICMYRHALPPGFVLK KDKKKEEKEDEISLEDLIERERS ALGPNVTKITLESFLAWKKRKR
18889	49257	Α	18998	1	322	
18890	49258	A	18999		822	MGKAKEEVFLADELVHAKTNR NKECDYSVTANSKIVVVTAGV RQQEGESRLNLVQRNVNVFKFI IPQIVKYSPDCIIIVVSNPVDILT YVTWKLSGLPKHRVIGSGCNL DSARFRYLMAEKLGIHPSSCHG WILGEHGDSSDHPGRPRPHPVPI KPPETLPGRDTSSWTSRGAHWR RNTQAAGRREHIDRHQQAIDR WNNVGFGRGGQRRARLLSDLT PTGKPSSYSVSLLAPPSAESYFH SIKPCIHSPS*CMQRPIEIKN/RD YSVTANSKIVVVTAGVRQQEG ESRLNLVQRNVNVFKFIIPQIVK YSPDCIIIVVSNPVDILTYVTWK LSGLPKHRVIGSGCNLDSARFR YLMAEKLGIHPSSCHGWILGEH GDSSDHPGRPRPHPVPIKPPETL PGRDTSSWTSRGAHWRRNTQA AGRREHIDRHQQAIDRWNNVG FGRGGQRRARLLSDLTPTGKPS SYSVSLLAPPSAESYFHSIKPCIH SPSPREIQFFGYTKARNPRVQK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18891	49259	Α	19000	1	1337	MEKPKRLRWLWGPAPSSLLAE
				1		PAPEETHAADFVFSARLLQRSP
		l				EPSLSWGGRALAEEQKAEGSSC
		1				GSAERLVAFAAKTRPYLSLCK
						MATLKEKLIAPVAEEEATVPNN
		1				KITVVGVGQVGMACAISILGKS
						RAMEHTTAGAPYTPRQQTCQC
						NMMSLADELALVDVLEDKLKG
		1				EMMDLQHGSLFLQTPKIVADK
		ŀ				DYSVTANSKIVVVTAGVRQQE
		ļ				GESRLNLVQRNVNVFKFIIPQIV
						KYSPDCIIIVVSNPRDI\LTYGTW
						KLSGLPKHRVIGSGCNLDSARF
						RYLMAEKLGIHPSSCHGWILGE
						HGDSSVAVWSGVNVAGVSLQE
						LNPEMGTDNDSENWKEVHKM
						VVESAYEVIKLKGYTNWAIGLS
			1			VADLIESMLKNLSRIHPVSTMV
						KGMYGIENEVFLSLPCILNARG
						LTSVINQKLKDDEVAQLKKSA
		ł				DTLWDIQKDLKDL
18892	49260	С	19001	462	728	
18893	49261	Α	19002	1	1033	MSGSFLMFLPKADMPAPMLVQ
		İ	ļ			PAERFDSWVWAGPEQLHSNQL
			ŀ			PGDADALLLGRNLTKGTRRLC
						TEEQVKIPQPEPGPPQKLCVLSA
		l		1		AGGQACLVLTFPRWTVWEEPP
						PPGLVGVKACARWWGKGEPD
		İ				RKAPAPPPPPPGTEKVDQTSE\E
		İ				RGTLIQRPPAPALPGWVRKTGR
ŧ		İ				P/PPPA/TGRLQQVPLLGASTQPE
						/PPGRGV/PPGVSPGLGEDVE/N
						DGMSLSYLAE*PQATSQRAP*G
		İ				ENTVALSGRLGRNSSSYTCRPA
						GERNFHPRGGPSSTAALHLRK
						MPGTPQCEGRRAVPRALPET*M
						LQGPGTTR*T/PPGPLAFSPPSSP
		ı	l			TGPQKVKSLGNSSARTILPPKW
18894						
	49262	Α	19003	2	212	
18895	49262 49263	A A	19003 19004	2	425	PTRPAAVAEDGGLKKCKISSYC
18895		₩		1		RSQPPARLISGEEHFSSKKCLA
18895		₩		1		RSQPPARLISGEEHFSSKKCLA WFYEYAGPDEVVGPEGMEKFC
18895		₩		1		RSQPPARLISGEEHFSSKKCLA WFYEYAGPDEVVGPEGMEKFC EDIGVEPEN\CDCTEKLQNKFDF
18895		₩		1		RSQPPARLISGEEHFSSKKCLA WFYEYAGPDEVVGPEGMEKFC EDIGVEPEN\CDCTEKLQNKFDF LRSQLNDISSFKNIYRYAFDFAR
18895		₩		1		RSQPPARLISGEEHFSSKKCLA WFYEYAGPDEVVGPEGMEKFC EDIGVEPEN\CDCTEKLQNKFDF

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18896	49264	Α	19005	229	1028	GASKRTGTPFPGFPAHPAREGR
				ļ		KMPVKKKRKSPGVAAAVAED
						GGLKKCKISSYCRSQPPARLISG
				ŀ		EEHFSSKKCLAWFYEYAGPDE
						VVGPEGMEKFCEDIGVEPENII
						MLVLAWKLEAESMGFFTKEEW
						VKG\MTSLQCNLHRKGYKPNF
						DFFAAHS*IDISSFKNIYRYAFDF
						ARDKDQR\SLDNDTAKS\MLAL
						LLGEGHGPLFS\VFYQVPGGQS
						KYRVMNKD\QWYNVSEFSRTV\
						HADLSNYDEDG\AWPVLLDEF
						VEWQKVRQTS
18897	49265	Α	19006	3	373	
18898	49266	Α	19007	3	418	GRAGSRKWLLLTGSLASTSPSL
		1				LSGQGPWAPLQRAMKPPGGES
		1	1			SNLF\GFPEEATPSSRPNRMSSNI
		:				FGPTEEPQNILKRTNPPGV*GSG
						IFDESTPVQTRQHLNPPG\GKTS
						DIFGSPVTATSRLAHPNKPKDH
	<u> </u>	<u> </u>			1604	VFLM
18899 18900	49267	A	19008 19009	3	684 81	
18901	49269	A	19010	1	465	
18902	49270	A	19010	120	344	
18903	49271	A	19012	3	1334	SPQLISSASRPAVRNSRHDATRP
1000	1/2/1	' '	1,,012			VVFQIMSGGSADYNSREHGGPE
		1				GMDP\DGVIESNWNEIVDNF/D
					1	VTMNLKESLL\RGI\YAYGFEKP
						SPIQQKA\IIPCIKGY*CGLLKLQ
						SGTGKTATF/SLISILQPVGRLSF
						KETQSTSYWAPTREL\AQQIQK
						VILALGDYMGATCHACIG\GTN
						VRNEMQKLQAEAPHIV\VGTPG
						RVFDMLNRRYLSPKW\IKMFVL
		1				DEADEMLSRGFKD\QIYEIFQKL
		1				NTSIQVVFASATMPTDVLEVTK
		1				KFMRDPIRILVKKEELTLEGIKQ
		1	1			FYINVEREEWKLDTLCDLYETL
						TITQAVIFLNTRRKVDWLTEKM
						HARDFTVSALHGDMDQKERDV
						IMREFRSGSSRVLITTDLLARGI
						DVQQVSLVINYDLPTNRENYIH
						RIGRGGRFGRKGVAINFVTEED
					-	KRFLRDIETFY\NTTVEEMPMN
18904	49272	A	19013	75	540	THE BREIDIT I WITT VEELVILLIVILLIVILLIVILLIVILLIVILLIVILLI
10704	17Z1Z		17013	1/3	שדע	L

SEQ ID	SEQ ID NO:	1	SEQ ID NO:	I.		Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		07/340,217	sequence	or peptide sequence	deterior, (possible nacicottae insertior)
18905	49273	IA	19014	774	1529	RWGERILLTWRKRQLQVPPSAL
						EAGGSSGLEDVLPLLQQADELH
		ļ				RGDEQGKREGFQLLL/TQQAW
		1				CMEAGRTFLWRLARAYSDMCE
		l				LTEEVSEKKSYALDGKEEAEAA
						LEKGDEE/SLTCHLWYAGALW
1		İ				SAG*AMRASRRRIQELALTFKE
						HCGQKPFALQP\KTPMAHFLLG
		l				RWCYQVSHLSWLEKKTATALL
		ŀ				ESPLSATVEDALQSFLKCYREL
						GKNSEARWWMKLALELPDVT
						KEDLAIQKDLEELEVILRD
18906	49274	Α	19015	97	452	GTERRTGKLPALLTPNLVQLRR
						LLFPGIQTI*VPAQGRGHLLQRP
						AAA/RPRTRSAPPAAAAAPPQV
						EARPP*LQPTRQEEAAELQTAR
					}	AGLWELWAGPQSPRDLNSLESI
						RITCSPTP
18907	49275	Α	19016	1299	1800	FWGGRSGTVGYGADMGKFKN
						NTTNNQS*K*YRNTSRNPDHKD
						*FLKGMDPKFPKNMHFA*KHN
						EKGLEKMQDNNMRAEAIKA/L
		1				PKPMEVQPRVPKSASRKLD/LA
						YTAHPKAGKHARAHIATGLRL
						CWPKAKAKDQTKAQASASAA
						APASVLAPAQASKGAQAPMKA
						LE
18908	49276	Α	19017	410	1323	RQEGTLIGVFIPAAAAAAAAVAA
						AAAAAAAVAAAAAAAAA
						AAAAAAAAAVAAAVAAAAA
						AV/GTSFSKPHKLMKEHKEKPS
						KDSREHKSAFKEPSRDHNKSSK
						ESSKKPKENKPLKEEKIVPKMA
						FKEPKPMSKEPKPDSNLLTITSG
						QDKKAPSKRPPISDSEELSAKK
						RKKSSSEALFKSFSSAPPLILTCS
						ADKKQIKDKSHVKMGKVKIES
						ETSEKKKSTLPPFDDIVDPNDSD
						VEENISSKSDSVPGQSPVTNIFF
						TSSSTPTILKSYQLDNSDQQGA
						QEASDKQIEGYFSGEVPAYRQG
		<u> </u>				ELILVASRQ

SEQ ID	-		SEQ ID NO:	4	§	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence	l	09/540,217	codon for peptide sequence	or pepude sequence	defetion, \=possible nucleotide insertion)
				sequence		
18909	49277	Α	19018	91	890	GAERLPEGPGKRMVFRRFVEV
1		l				GRVAY\VSFGPHARK/MWVAIV
		l				DVF*SEQGLLVE*PCTQVR\RQA
		1		1		\MPFRCMQLN*FHPSRFPNSAQP
						ESMFRQAWQK\ADI\NTKWGSP
				:		HDGAKKIEAQRKGKPKDDQIF
		1				D\RFKVMEGQKKMREQE*SKE*
		1				S*RSFKRAALLEKLSPKKKHLG
		l				T*GYCCLLLLLAAAAAAAAAA
						AACCLLACCLKVPSKKRSPAAS
		1				KKAPAQKVPAQK\ATG\QKAAP
		1				APKAQKGQKAPAQKAPAPKAS
		1				GQESISGNHKK
18910	49278	Α	19019	280	1249	SLNEGIMYAQP\VTNTKEVK\W
		1				QKV\LYERQPFSLDKLWWDRRF
		1				LGRSSGKNIHAPENTQY\WGCG
				:		*FESSV\VIQQLCSVCVFVVIR\W
						YMDEGLLAPHWLVGTGLASSL
1	,					IGYVLFDLIDGGEGRKKSG\QTR
		ŀ				WADL*ELP*SFITFTYG\FSPVLK
						TLTEFCSALTPIYAMSVFMLLR
		İ				HLIFFDYG\ANAAIV\SST\LLVN
		ł				MAIFAS\VCWASRLP\RS\LHAFI
						MVA\FAIQIFCPGGPML\QKKLK
						\ACTP\RS\YVGVTL\LFAFSAVG\
						GLLSISAVGA\VLLA\LLLMSISC
						LC\PFYLHFACSLFKENI/HMGP
						WDEA\EI\KEDLSPGSLS
18911	49279	Α	19020	1	310	
18912	49280	Α	19021	2	411	
18913	49281	Α	19022	374	1275	VLTRLIRKRERRKAQSQRERCR
		1				CVLAKDFLAGGVTAAISKMAV
		İ				APTE/RVKLLLQVQSASKQITAD
		1				KQYTGVVDCMVRIPKE/QGVLS
		l				LWH\GIQFWHKFAGSLASGGAP
		1				GATSLCFVYPL\DFARTRLAAD
		ŀ				VGKAGAEREFRGLG*PAGLRIY
		1				KSDGK/LRGLYQGF\NVSVQGIII
		İ				YRAAYFGIYD\TAKG\MLPDP\Q
		ł				NTHIVISWMIAQTVTAVARVDF
		ļ				PIPFDT\VRRRNE*MQSRAPKEL
						TIHVHQARFDCWAEGLLRDEG
						G\KAFFKG\AWVPMFLRGHGVG
10011	10000	_	10000	161	705	AFCALSLYGWKSKEGTHKLIS
18914	49282	Α	19023	461	705	TERKKKGILPGNGNGFMSLRRT
}						IRVWRNTPNYLGTNAVSP\RQD
1						*IGKKGVCQRKQAGPCPPSNIQ
10015	40202	-	10024		400	GILEWRKHRISETKP
18915	49283	Α	19024	2	408	

LGRRKQLEDYLTKILKMPMYR NYHATTEFLDISQLSFIHDLGPK GIEGMIMKR 18919 49287 B 19028 37 147	SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
18916	NO:		hod	1	1		
18916 49284 A 19025 1 885 MTQHAGKRDSELPVIVQDTET		sequence		09/540,217	1	of peptide sequence	deletion, \=possible nucleotide insertion)
VPSVMDGKEYQAGRFARGLRL QCFRRRRRRTRRRRKEEEE EEEEEEEEEEEVEEGMKKEE EGRRKKKKKKAEEEEKKEEE EGRRRKKKKKKAEEEEKKK KEEEEKKKKKRRREEPPRPRNDDD ARRRRKEEEEGRRSM/LGKRD SELPVIVQDTETVPSVMDGKEY QAGRFARGLRLQCFRRRRR RTRRRRKEEEEEEEEEEEEEE EVEEEGMKKEEEGRRKKKK KKAEEEEEKKKKEEEKKRR REEPPRPRNDDDARRRKEEEE GRRSMKKEEEEEEEEEEE GRRSMKKEEEEEEEEEEE EEEEEEEEKKRRIGPASKMM GGFEELRRGLNVGPTMAIFFLQ SAALAPAILVFCPSLKSNISKON FQFIMFCQLLGRVTIPFLVSQRS RIEVGRHSLSTLNAETAHELYL GRTEPDECGKTAGLGTGLDLK 18917 49285 A 19026 1 378 18918 49286 A 19027 36 3415 ESAANAQVLAAPSPCSPFAFTL SKVNMSLKNEPVNTSALQKIA ADMSNIIENLDTRELHFEGEEV DYDVSPSDPKIQEVYIPFSAIYN TQGFKEPNIQTYLSGCPIKAQVL EVERFTSTTRVPSINLYTIELTH GFEK WQVKRKFKHFQEFHREL LKYKAFIRIPPTRRHTFRRQNV REEPREMPSLPRSSENMIREEQF LGRRKQLEDYLTKILKMPMYR NYHATTEFLDISQLSFIHDLGPK GIEGMIMKR	1				sequence		
VPSVMDGKEYQAGRFARGLRL QCFRRRRRRTRRRRKEEEE EEEEEEEEEEEVEEGMKKEE EGRRKKKKKKAEEEEKKEEE EGRRRKKKKKKAEEEEKKK KEEEEKKKKKRRREEPPRPRNDDD ARRRRKEEEEGRRSM/LGKRD SELPVIVQDTETVPSVMDGKEY QAGRFARGLRLQCFRRRRR RTRRRRKEEEEEEEEEEEEEE EVEEEGMKKEEEGRRKKKK KKAEEEEEKKKKEEEKKRR REEPPRPRNDDDARRRKEEEE GRRSMKKEEEEEEEEEEE GRRSMKKEEEEEEEEEEE EEEEEEEEKKRRIGPASKMM GGFEELRRGLNVGPTMAIFFLQ SAALAPAILVFCPSLKSNISKON FQFIMFCQLLGRVTIPFLVSQRS RIEVGRHSLSTLNAETAHELYL GRTEPDECGKTAGLGTGLDLK 18917 49285 A 19026 1 378 18918 49286 A 19027 36 3415 ESAANAQVLAAPSPCSPFAFTL SKVNMSLKNEPVNTSALQKIA ADMSNIIENLDTRELHFEGEEV DYDVSPSDPKIQEVYIPFSAIYN TQGFKEPNIQTYLSGCPIKAQVL EVERFTSTTRVPSINLYTIELTH GFEK WQVKRKFKHFQEFHREL LKYKAFIRIPPTRRHTFRRQNV REEPREMPSLPRSSENMIREEQF LGRRKQLEDYLTKILKMPMYR NYHATTEFLDISQLSFIHDLGPK GIEGMIMKR	18916	19284	Δ	19025	<u>1</u> 1	885	MTOHAGKROSEI PVIVODTET
QCFRRRRRRTRRRRKEEEE EEEEEEEEVEEGMKKEE EGRRKKKKKAEEEEKKK KEEEEKKRRREEPPRPRNDDD ARRRKEEEEGRRRSM/LGKRD SELPVIVQDTETVPSVMDGKEY QAGRFARGLRLQCFRRRRRR RTRRRRKEEEEEEEEEE EVEEEGMKKEEEGRRKKKKK KKAEEEEEKKKEEEEEEEEE EVEEEGMKKEEEGRRRKKKKK KKAEEEEEKKKREEEEEEEE GRRRSMKKEEEEEEEEE EVEEEGMKKEEEGRRRKKKK KKAAEEEEEKKKREEEEEEEE GRRRSMKKEEEEEEEEE GRRRSMKKEEEEEEEEE GRRRSMKKEEEEEEEEE GRRRSMKKEEEEEEEEE GRRRSMKKEEEEEEEEE GRRRSMKKEEEEEEEEE GRRRSMKKEEEEEEEEE GRRRSMKKEEEEGEEEE GRRRSMKKEEEEGEEEEE GRRRSMKKEEEEGEEEEE GRRRSMKKEEEEGEEEEE GRRRSMKKEEEEGEEEEE GRRRSMKKEEEEGEEEEE GRRRSMKKEEEEGEEEEE GRRRSMKKEEEEGEEEEE GRRRSMKKEEEEGEEEEE GRRRSMKKEEEEGEEEEE GRRRSMKKEEEEGEEEEE GRRRSMKKEEEEGEEEEE GRRRSMKKEEEEGEEEEE GRRRSMKKEEEEGEEEEE GRRRSMKKEEEEGEEEEEE EEGEEKKRRR REPPPRPNDDDARRKKEEE GRRSMKKEEEEGEEEEE GRRSMKKEEEEGEEEEE GRRSMKKEEEEGEEEEE GRRSMKKEEEEGEEEEE GRRSMKKEEEEGEEEEE GRRSMKKEEEEGEEEEE GRRSMKKEEEEGEEEEE GRRSMKKEEEEGEEEEEEE GRRSMKKEEEEGEEEEE GVEEGMKKKE KKAEEEEKKKREEE GRRSMKKEEEEGEEEEEEE GVEEGMKKKE KKAEEEEKKKKEEEE GRRSMKKEEEEGEEEEEE GVEEGMKKEE GEEMKKREEEE GVEEGMKKKE KKAEEEEKKKKEEEE GRRSMKKEEEEGE GRRSMKKEEEEEEEEEE GVEEGMKKEE GEGEMKKE KKAAEEEEEKKKKEEEEEEEEEEEE GVEEGMKKEE GEEMKKREEEE GVEEGMKKKEEE GVEEGMKKKEEEE GVEEGMKKEEE GVEEGMKKKEEEE GVEEGMKKEE GEGRKSMKKEEEEEEEEEEEEE GVEEGMKKEE GEGEGKKKKEEEEEEEEEEEEEEE GVEEGMKKKEEEEEEEEEEEEEEEEEEEEEEE GVEEGMKKEEEEEEEEEEEEEEEEEEEEEEEEEEEEEEEE	10010	17204		17023		1003	, ,
EEEEEEEEEEVEEGMKKEE EGRRRKKKKKKAEEEEKKK KEEEKKRRREEPPRPNDDD ARRRRKEEEGRRSM/LGKRD SELPVIVQDTETYPSVMDGKEY QAGRFARGLRQCFRRRRRR RTRRRRKEEEEEEEEEEEEEEEEEEEEEEEEEE							,
EGRRRKKKKKKAEEEEEKKK KEEEEKKRRREEPPRPNDDD ARRRKEEEGRRSM/LGKRD SELPVIVQDTETVPSVMDGKEY QAGRFARGIRLQCFRRRRRR RTRRRRKEEEEEEEEEEEEEEEEEEEEEEEEEEEEEEEE							l '
KEEEKKRRREEPPRPRNDDD ARRRKEEEGRRRSM/LGKRD SELPVIVQDTETVPSVMDGKEY QAGRFARGLRLQCFRRRRRR RTRRRKEEEEEEEEEEEEEEEEEEEEEEEEEEE							
ARRRKEEEEGRRRSM/LGKRD SELPVIVQDTETVPSVMDGKEY QAGRFARGLRLQCFRRRRRR RTRRRRKEEEEEEEEEEEEEEEEEEEEEEEEEE							
SELPVIVQDTETVPSVMDGKEY QAGRFARGLRLQCFRRRRRR RTRRRRKEEEEEEEEEEEEEEEEEEEEEEEEEE							
QAGRFARGLRLQCFRRRRRR RTRRRRKEEEEEEEEEEEEEEEEEEEEEEEEEE				l i			
RTRRRRKEEEEEEEEEEEEEEEEEEEEEEEEEEEEEEEE							
EVEEEGMKKEEEGRRRKKKK KKAEEEEKKKRR REEPPRPRNDDDARRRRKEEEE GRRRSMKKEEEEEEEEEEEEE EEEEEEEEEKKKRRIGPASKMM GGFEELRRGLNVGPTMAIFFLQ SAALAPAILYFCPSLKSNISKCN FQFLMFCQLLGRVTIPFLVSQRS RIEVGRHSLSTLNAETAHELYL GRTEPDECGKTAGLGTGLDLK 18917 49285 A 19026 1 378 18918 49286 A 19027 36 3415 ESAANAQVLAAPSPCSPFAFTL SKVNMSLKNEPRVNTSALQKIA ADMSNIIENLDTRELHFEGEEV DYDVSPSDPKIQEVYIPFSAIYN TQGFKEPNIQTYLSGCPIKAQVL EVERFTSTTRVPSINLYTIELTH GEFKWQVKRKFKHFQEFHREL LKYKAFIRIPIPTRHTFRRQNV REEPREMPSLPRSSENMIREEQF LGRRKQLEDYLTKILKMPMYR NYHATTEFLDISQLSFIHDLGPK GIEGMIMKR							
KKAEEEEKKKKEEEKKRRR REEPPRPRNDDDARRRRKEEEE GRRRSMKKEEEEEEEEEEEEE EEEEEEEEEKKKRIGPASKMM GGFEELRRGLNVGPTMAIFFLQ SAALAPAILYFCPSLKSNISKCN FQFLMFCQLLGRVTIPFLVSQRS RIEVGRHSLSTLNAETAHELYL GRTEPDECGKTAGLGTGLDLK 18917 49285 A 19026 1 378 18918 49286 A 19027 36 3415 ESAANAQVLAAPSPCSPFAFTL SKVNMSLKNEPRVNTSALQKIA ADMSNIIENLDTRELHFEGEEV DYDVSPSDPKIQEVYIPFSAIYN TQGFKEPNIQTYLSGCPIKAQVL EVERTSTTRVPSINLYTIELTH GEFKWQVKRKFKHFQEFHREL LKYKAFIRIPIPTRHTFRRQNV REEPREMPSLPRSSENMIREEQF LGRRKQLEDYLTKILKMPMYR NYHATTEFLDISQLSFIHDLGPK GIEGMIMKR							
REEPPRPRNDDDARRRRKEEEE GRRRSMKKEEEEEEEEEEEE EEEEEEEEEKKKRRIGPASKMM GGFEELRRGLNVGPTMAIFFLQ SAALAPAILYFCPSLKSNISKCN FQFLMFCQLLGRVTIPFLVSQRS RIEVGRHSLSTLNAETAHELYL GRTEPDECGKTAGLGTGLDLK 18917 49285 A 19026 1 378 18918 49286 A 19027 36 3415 ESAANAQVLAAPSPCSPFAFTL SKVNMSLKNEPRVNTSALQKIA ADMSNIIENLDTRELHFEGEEV DYDVSPSDPKIQEVYIPFSAIYN TQGFKEPNIQTYLSGCPIKAQVL EVERFTSTTRVPSINLYTIELTH GEFK WQVKRKFKHFQEFHREL LKYKAFIRIPIPTRRHTFRRQNV REEPREMPSLPRSSENMIREEQF LGRRKQLEDYLTKILKMPMYR NYHATTEFLDISQLSFIHDLGPK GIEGMIMKR							
GRRRSMKKEEEEEEEEEEEEEEEEEEEEEEEEEEEEEEEEE							
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FQFLMFCQLLGRVTIPFLVSQRS RIEVGRHSLSTLNAETAHELYL GRTEPDECGKTAGLGTGLDLK 18917 49285 A 19026 1 378 18918 49286 A 19027 36 3415 ESAANAQVLAAPSPCSPFAFTL SKVNMSLKNEPRVNTSALQKIA ADMSNIIENLDTRELHFEGEEV DYDVSPSDPKIQEVYIPFSAIYN TQGFKEPNIQTYLSGCPIKAQVL EVERFTSTTRVPSINLYTIELTH GEFK WQVKRKFKHFQEFHREL LKYKAFIRIPITRRHTFRRQNV REEPREMPSLPRSSENMIREEQF LGRRKQLEDYLTKILKMPMYR NYHATTEFLDISQLSFIHDLGPK GIEGMIMKR			ļ				•
RIEVGRHSLSTLNAETAHELYL GRTEPDECGKTAGLGTGLDLK 18917 49285 A 19026 1 378 18918 49286 A 19027 36 3415 ESAANAQVLAAPSPCSPFAFTL SKVNMSLKNEPRVNTSALQKIA ADMSNIIENLDTRELHFEGEEV DYDVSPSDPKIQEVYIPFSAIYN TQGFKEPNIQTYLSGCPIKAQVL EVERFTSTTRVPSINLYTIELTH GEFK WQVKRKFKHFQEFHREL LKYKAFIRIPIPTRRHTFRQNV REEPREMPSLPRSSENMIREEQF LGRRKQLEDYLTKILKMPMYR NYHATTEFLDISQLSFIHDLGPK GIEGMIMKR 18919 49287 B 19028 37 147	1			 			ł l
GRTEPDECGKTAGLGTGLDLK 18917 49285 A 19026 1 378 18918 49286 A 19027 36 3415 ESAANAQVLAAPSPCSPFAFTL SKVNMSLKNEPRVNTSALQKIA ADMSNIIENLDTRELHFEGEEV DYDVSPSDPKIQEVYIPFSAIYN TQGFKEPNIQTYLSGCPIKAQVL EVERFTSTTRVPSINLYTIELTH GEFKWQVKRKFKHFQEFHREL LKYKAFIRIPIPTRRHTFRRQNV REEPREMPSLPRSSENMIREEQF LGRRKQLEDYLTKILKMPMYR NYHATTEFLDISQLSFIHDLGPK GIEGMIMKR 18919 49287 B 19028 37 147							
18917 49285 A 19026 1 378	ļ			!			1
18918 49286 A 19027 36 3415 ESAANAQVLAAPSPCSPFAFTL SKVNMSLKNEPRVNTSALQKIA ADMSNIIENLDTRELHFEGEEV DYDVSPSDPKIQEVYIPFSAIYN TQGFKEPNIQTYLSGCPIKAQVL EVERFTSTTRVPSINLYTIELTH GEFKWQVKRKFKHFQEFHREL LKYKAFIRIPIPTRRHTFRRQNV REEPREMPSLPRSSENMIREEQF LGRRKQLEDYLTKILKMPMYR NYHATTEFLDISQLSFIHDLGPK GIEGMIMKR							GRTEPDECGKTAGLGTGLDLK
SKVNMSLKNEPRVNTSALQKIA ADMSNIIENLDTRELHFEGEEV DYDVSPSDPKIQEVYIPFSAIYN TQGFKEPNIQTYLSGCPIKAQVL EVERFTSTTRVPSINLYTIELTH GEFKWQVKRKFKHFQEFHREL LKYKAFIRIPIPTRRHTFRRQNV REEPREMPSLPRSSENMIREEQF LGRRKQLEDYLTKILKMPMYR NYHATTEFLDISQLSFIHDLGPK GIEGMIMKR			Α		ļ		
ADMSNIIENLDTRELHFEGEEV DYDVSPSDPKIQEVYIPFSAIYN TQGFKEPNIQTYLSGCPIKAQVL EVERFTSTTRVPSINLYTIELTH GEFKWQVKRKFKHFQEFHREL LKYKAFIRIPIPTRRHTFRRQNV REEPREMPSLPRSSENMIREEQF LGRRKQLEDYLTKILKMPMYR NYHATTEFLDISQLSFIHDLGPK GIEGMIMKR 18919 49287 B 19028 37 147	18918	49286	Α	19027	36	3415	`
DYDVSPSDPKIQEVYIPFSAIYN TQGFKEPNIQTYLSGCPIKAQVL EVERFTSTTRVPSINLYTIELTH GEFKWQVKRKFKHFQEFHREL LKYKAFIRIPIPTRRHTFRRQNV REEPREMPSLPRSSENMIREEQF LGRRKQLEDYLTKILKMPMYR NYHATTEFLDISQLSFIHDLGPK GIEGMIMKR 18919 49287 B 19028 37 147	1						1
TQGFKEPNIQTYLSGCPIKAQVL EVERFTSTTRVPSINLYTIELTH GEFKWQVKRKFKHFQEFHREL LKYKAFIRIPIPTRRHTFRRQNV REEPREMPSLPRSSENMIREEQF LGRRKQLEDYLTKILKMPMYR NYHATTEFLDISQLSFIHDLGPK GIEGMIMKR 18919 49287 B 19028 37 147							1
EVERFTSTTRVPSINLYTIELTH GEFKWQVKRKFKHFQEFHREL LKYKAFIRIPIPTRRHTFRRQNV REEPREMPSLPRSSENMIREEQF LGRRKQLEDYLTKILKMPMYR NYHATTEFLDISQLSFIHDLGPK GIEGMIMKR 18919 49287 B 19028 37 147							1
GEFKWQVKRKFKHFQEFHREL LKYKAFIRIPIPTRRHTFRRQNV REEPREMPSLPRSSENMIREEQF LGRRKQLEDYLTKILKMPMYR NYHATTEFLDISQLSFIHDLGPK GIEGMIMKR 18919 49287 B 19028 37 147	l						
LKYKAFIRIPIPTRRHTFRRQNV REEPREMPSLPRSSENMIREEQF LGRRKQLEDYLTKILKMPMYR NYHATTEFLDISQLSFIHDLGPK GIEGMIMKR 18919 49287 B 19028 37 147							EVERFTSTTRVPSINLYTIELTH
REEPREMPSLPRSSENMIREEQF LGRRKQLEDYLTKILKMPMYR NYHATTEFLDISQLSFIHDLGPK GIEGMIMKR							GEFKWQVKRKFKHFQEFHREL
LGRRKQLEDYLTKILKMPMYR NYHATTEFLDISQLSFIHDLGPK GIEGMIMKR 18919 49287 B 19028 37 147							LKYKAFIRIPIPTRRHTFRRQNV
NYHATTEFLDISQLSFIHDLGPK GIEGMIMKR							REEPREMPSLPRSSENMIREEQF
GIEGMIMKR 18919 49287 B 19028 37 147							LGRRKQLEDYLTKILKMPMYR
18919 49287 B 19028 37 147							NYHATTEFLDISQLSFIHDLGPK
							GIEGMIMKR
	18919	49287	В	19028	37	147	
18920 49288 B 19029 10 264	18920	49288	В	19029	10	264	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18921	49289	Α	19030	3	1255	HASGGRAANMAADRGAVQQQ
						SQDMMEVDRRVESEESGDEEG
						KKHSSGIVADLSEQSLKDGEEP
	İ					G\EEDPEEEHELPVDMETINL\D
						RDAEDF\DLNHYRIGKIEGFEVL
						EES*RLSCLRQNLI\KCL\ENLGR
						ELPESFRELDLYDNQIKKIENLE
						ALTELEILDISFNLLRNIEGVDK
						LTRLKKLFLVNNKISKIENL\SN
					:	LHQL\RMLGAGDLNRIRGQFSK
						YPTTL\TNL\ESWFLGKKTKITN
						FQNPGCASPNLTVLSMQSNRLT
						KIEGLQN\LVNLRELYFSHNGIE
						VIEGLENN\NKLT\MLDIASNRIK
						KI*KISAHLTEL\QEFWDGTTIPL
						KSWSDLDELEGEPGALEDSVP
						WGGNPLARRTPKYRRK\VMLA\
						LPSVRQDRCPRSVRFLKSFLGSL
						MWSLSSEELPSHGVFNPPCCS
18922	49290	A	19031	138	564	YREVMVSES*ETPAGARGRPYY
10922	49290	A	19031	136	304	FSAPGTAP\PAINVHPPPPSLSAT
	ļ	1				PHPPQPQPPPPHQHNAKARVAT
		1				IRTKRTSNCRIRSRKVRKSPPEK
						WVGFNRRPKASCPSPPGAARV
		1				
						DVGGETERREQAAAPGEMGK
10022	40201	ļ. —	10022	1404	1506	WARPGEEYFHS
18923	49291	Α	19032	1404	1586	ENESRFSDRNQASAGLGYLSDS
			}			L*QWIVGNGHATDLWQNCSTS
10024	10000	ļ. —	10000	107	701	SSGNVHHCFSSSPNGSG
18924	49292	A	19033	187	701	AELAARMLLLLLSIIVLHVAAL
						VLLFVSTIVSQWIVGNGHATDL
						WQNCSTSSSGN\VHHCFSSSPNE
						W\LQSCSRGTMDPVDSSFSILSL
		İ				FLFFCQLFTLTKGG\RFYITGIFQ
						ILAGLCV\MSAAAIYTVRHPEW\
						NLNSGY\S*RFA*I\LAWVAFPL\
					==-	ALLSGVIYVILRKRE
18925	49293	A	19034	1	493	
18926	49294	Α	19035	105	307	
18927	49295	A	19036	32	302	
18928	49296	В	19037	119	641	
18929	49297	A	19038	496	653	
18930	49298	C	19039	568	747	
18931	49299	Α	19040	522	656	ESSQSHGKNFYMGKRIKLSLY
		1				WNVWRMLNSKSETTNCPLIRH
						RLSKRVGAPFLPAVLVLTKITIY
	<u> </u>	L				QT*KFQIRNHKLPSNKTSAI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18932	49300	Α	19041	1111	277	RHVVAVVLGDVGRSPRMQYH
						ALSLAMHGFSVTLLGFCNSKPH
			E			DELLQNNRIQIVGLTELQSLAV
						GPRVFQYGVKVVLQAMYLLW
		İ				KLMWREPGAYIFLQNPPGLPSI
			:			AVCWFVGCLCGSKLVIDWHNY
						GYSIMGLVHGPNHPLVLLAKW
		İ				YEKFFGRLSHLNLCVTNAMRE
						DLADNWHIRAVTVYDKPASFF
						KETPLD\RTHRLFTKLTNMHSPF
						RARSEPEDPVTERSAFTERDAG
						SGLVTRLRERPALLVSST\SWTE
						DEDFSILLAALESRV*TTDS/SIG
						HNLPSLVCVITGKGPPREYYSR
				İ		LIHQKHFQHIQVCTPWLEAEDY
						PR/ILGSVDLGVCLHTSCSGLDL
		İ				PMKVVDMFGCCLPVCAVNFK
		İ		ļ		WQEQNPNLSGDSFTDPPLRRKQ
	ļ					CRASCPQ*GPAPGLHELVKHEE
						NGLVFEDSEELAAQLQVLFSNF
						PDPAGKLNQFRKNLRESQQLR
						WD*SWGSATPNPMMSSCRTTE
18933	49301	В	19042	110	2294	
18934	49302	В	19043	81	1538	
18935	49303	Α	19044	21	1158	AVVAALPTSSSSRHSLTQKPGS
						RRLRIESLLPPSSCEIFLIFSYIFR
						LCEKPYHQTRSASNVTNKTDPR
			<u> </u>			SMNS\RVFIGNLNTLV\VKKSDV
						EAIFSK\YGKI\VGCSVHKGLCLS
						VQVMLMRRNAPGLL*QGE\DG
						RMI/ALGQVLDINLAAEPK\VN\
						REKSSVKRSAAEMYG/SQ*QNT
						LLRPLYFSSSFDLDYDFQRDYY
					1	DRMYSYPARVPPPPPIARAVVP
						SKRQRVSGNTS\RSGPRGFNSKE
						WNSGVSSKFWKR*KGDDLQ\AI
						KEELT\QINQK\VDSLLENLEKIE
						KATGANKQLEMK\ND\KSEEEQ
						SSSSREGKMRTNVKDGV*GGV
						PDDSAEGGGPTWMNDDNVKS
				†		GGMTSWELI\KDDEKEAEE\GE
						DD\RD\SANGGG
10000					•	
18936	49304	В	19045	93	303	
18936 18937	49304 49305	B A	19045 19046	93 1	303 246	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18939	49307	Α	19048	584	1664	SGLFLFLFFYSEIFLIFSYIFRLCE
						KPYHQTRSASNVTNKTDPRSM
						NS\RVFIGNLNTLV\VKKSDVEA
						IFSK\YGKIVG\CSCFIKGFAFPFK
						*C**EEMPRACCSGRRVGRIDLL
						GQVLDI*PGLAEPK\VNPREKPG
		İ		:		VETILPAGDVRGQ*QNTLLRPL
						YFSSSFDLDYDFQRDYYDRMY
						SYPARVPPPPPIARAVVPSKRQR
						VSGNTSRRGK\SGFNSKEWNSG
						VSSKFWKR*KGDDLQ\AIKEEL
						T\QINQK\VDSLLENLEKIEKAT
						GANKQLEMK\NDK\SEEEQSSSS
						REGKMRTNVKDGV*GGVPDDS
						AEEGGPTWMNDDNVKSGGMT
						SWELI\KDD\EKEAEE\GEDDKR
						QGPMARNDS
18940	49308	Α	19049	158	1001	KRLGSSAISMSKYKLIMLR\HGE
						GAWNKENRFCSWVDQ\KLNSE
						GMEEARNCGKQLKALNFEFDL
						VFTSVLNRAIPTARL\IL\EELGQ
						EWVPVESSWRLNERH\YGAL\I
						GLNREQ\MA\LNHGE\EQVRL\W
	1					RRSYNVTP\PPIEE\SHPY\YQEIY
						NRPGGIKVCDVPL\DQLPRSES\
						LKDVL\ERFLPYW\NERIAPEV\L
						RG\KTILISAHGK*AVGALLKHL
						GRYPSGLKTSINIYSFLLGVPILL
						\ELD\ENLRAVGPH\QFLGDQEAI
						QAAIKKVEDQGKVKQAKK
18941	49309	A	19050	1	521	SASWVGVAMASRVLSAYVSRL
						PARFAP*PRVRMLAVARH\LST
		ŀ				ALCSEGTQTRLGTLQPALVLA\
						QVPW*ELHMLCRQYSDMP/PRL
						TLEGIQ\DRVRLA*LKTLSDKVD
						PREAFQ*ILHFMKDLGLDSLDP
						KWKII\MALED\EFGFEIP\DI\DA
10040	10010	ļ	10071			EKLM\TPPRKLVDLHCQDKKD
18942	49310	A	19051	1	516	IRALTMIWRRALLAGTRLVWS
		İ				RSGSAGWLDRAAG\LR\DCG\TA
						ASGMESNTSSSLENLATAPVNQ
						I\QETISDNCVVIFSKTSCSYCTM
						AKK\LFHDM\NVN\YKVVELDL
						LEYGN\QFQDASLQK*LVERTV
						PRIFVNGTFIGGATDTHRLHKE
10042	10211	_	10052	025	1000	GKLLPLVHQCYLKK\SKRKEFQ
18943	49311	Α	19052	825	1080	KRKIHQNFHKLSLKKAIEKDYS
						ESGSSVNQYKRIRPTDNVVIKQ
						Y\WPGAVAHTCNSSTSVGQGR
10044	40212		10053	80	120	WIT*GQEFETSLANVVKPHLY
18944	49312	Α	19053	80	429	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18945	49313	Α	19054	3	752	FKNAVGNLAMEGGGGIPLETL
						KEESQSRHVLPASFEVNSLQKS
						NWGFLLTGLVGGTLVAVYAVA
						TPF\VTPALRKVCLPFVPATTK\
		Ì				QIENVVKMLRCRRRDPLWDIGS
						GGRDGIVHRRLRRKGFTAVGF
						MEFKPMG*FWVFPRY\RAWRE
}						GVHGSAKFYISDLWKVTFSQYS
						NVVIFGVPQMMLQLEKKLERE
						LEDDARVIACRFPFPHWTPDHV
						TGKGIDTVWAYDASTFRGREK
						RPCTSMHFQLPIQA
18946	49314	Α	19055	157	829	TWGKGDPKKPRGKMSSYAFFV
						QTCRGG\HKKKHPDASVNFSEF\
						SKKCSERWKTMSA*R/EKGKFE
						DMAKA\DKARY\EREMKTYIPP
ļ						QRGRQKRKFKDSQLHPRRPPSG
						LLSSSCSEYRPKIK\GEHP\GL\SI
						GDVAKKLGRDVGINTAAD\DK
						QP\YEKK\AAKLKEKYEKDIA\A
						YRA\KGKPDAAKKG\VVKAEKS
					İ	KKKKEEEEDEEDEEDEEEEDE
						EDEEDEEDDG
18947	49315	Α	19056	85	267	GHLSHAWCGTYKPWPSPKEQS
						RAWKKDGLLFPPPP/PLFPTGLK
						VRGRESF*SVELIRPQG
18948	49316	Α	19057	1	1158	
18949	49317	Α	19058	452	517	
18950	49318	Α	19059	93	1069	YEGVTTSRLPELPRGCLVLQEQ
						ELVQMSGMEATVTIPIWQNKP
						HGC/SLRSVVRRIGTNLPLKPCA
						RASFETLPNISDLCLKRCAPSSL
					1	PWL\AWLGFLAE*KGSQKARV\
:						RSDTRPLRHTWKPSPLIVMQRN
						ASVPNLRGSEERLLALKKPA/L
						ASPKAALLSCRTS*ATLRTKICK
		1				DSGKLMQLRLH*RQISYLQEVQ
		İ				MSLLPLPCFGSSFHSTTSFCH**/
						PSPRRQRWRSLSFHQSPCFVLP
		1				ALNVANQNTKLPAVRLKRMTA
		1				SLCPRPAALQDMMGILKGLFTG
		1				MKQSQDLNRSLLKEEDPAVLIS
L	<u> </u>	1]		<u> </u>	EVLRRKFALKEEDISRKGN

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18951	49319	Α	19060	162	1410	GQASCPPPALGQQTPASSYTVQ
						IQTYAPSSPHPRQGPGQGSSDPR
						GLRGAIPRGRQGQGV*QQGPRC
						RPPSSPVLPHIPQLSCCGHDGRD
į						PEEGLGILELTLLHHAHPWSPN
						SW*KDS\PSPHGEFPAHSEPHSR
						DQTSPMATTKHEVQPLPQASLS
		İ				QPSLGIGCLVDWLRQLRTTQGS
						/PQDAAQDSTWERP*DAS/GKP
						WLL*PGVPTPGPPCQAKERKPP
						FMGNCPL*SRVPWPSRLPEEL/P
		İ				AMPLAPPVAGEAAGPASSHQSL
						SPSQDPTPLPGADRREKERGGE
1						EGSSWPQTPRLTTHTTSPPPEKQ
						QQIESPPIRLSSFKTGFHLASSGH
						SYAIKPNSKTKAEKVGQVRVPR
		l				SHTIREQRQGKRPGEHHRRTER
ł						DAKPQGEEDHQATVVSQKLGT
		l				DRRLIKKQRTSVLSATAES
18952	49320	Α	19061	1	340	GSVGSGTTLNQLGFWSSPGTSP
		1				LEPPCPWLDSHIFPVPFWKDVA
						FGEGVCLPPGAPQPRRICPRRG
						QQPEFQQEEPETIPERTPADPGV
						LPPAFPGICLPRRLRAPPDPGPA
						A*AVPPVADA*GSVGSGTTLNQ
						LGFWSSPGTSPLEPPCPWLDSHI
		ŀ				FPVPFWKDVAFGEGVCLPPGAP
						QPRRICPRRGQQPEFQQEEPETI
		1				PERTPADPGVLPPAFPGICLPRR
		1				LRAPPDPGPAA
18953	49321	Α	19062	3	396	IPLASLHLALPGHQRGPPPGRLP
		İ				SCSSQGTVPPPRAGPYQAGCPG
						TAPAPAAPASAPWSAPQASATG
						GTAHAAGPGSGGARSLPPARA
						HPPARCWQGRPLRSLLL*PSSVP
						ETGGPEKARLVKALSPNTAP
18954	49322	Α	19063	563	1269	LEESTGRITHFKATLHCTMSNN
						LAWGGTKWLQESRVFFHIRRLP
				1		SDASAGPGCVYRVVPDGGSQG
		1				RPQGAPP*AAQA/GTPRGRPP\P
				1		GAPSALLPPPPSGTSVGPGGLQ
						QRRPPTGQEDSPTEPGFWKPRL
						WRSHLATSLATRCTMQGEPAR
						AHPPGGRGPAASEERHQRGPPP
					1	GGCFAALRYCATPRPDLSGRCP
						GTAPAPAAPASAPWTRLRHQPL
		L		<u> </u>		AAQLMQPGQVWWGPAASW

sequence 09/540,217 codon for peptide of peptide sequence deletion	codon, /=possible nucleotide n, \=possible nucleotide insertion)
QLEE KFRE RPNL	VALVADEL DEECDOL DODCD
QLEE KFRE RPNL	ZAVADEL DEECDEL DEDCE
KFRE RPNL	KAKARELREECRSLRSRCD
RPNL	RVSVMENEMNEMKREG
1 1 1 1 1 1	KRIKRNEQSLQEIWDYVK
	HLIGVPESDGENGTKLEN
1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	DIIQENFPNLARQANIQIQEI
1 1 1 1 1	PORYSSRRATPRHIIVRITK
	KEKMLRAAREKGRVPHK
	RLTADLSSETLQARREWG
	LKRIFSFPQRGGTGGRRM
	TAELIEQAAQYTSAVRDRE RGYKIPVIENLGATLDQFD
	SDNEIRKLDGFPLLRRLKT
1 !	I/NNRIC/RVLDFQKVK/LK
1 1 1 1 1	EAEKMFKGKRGAQLAKDI
1 1 1 1 1 1 1	SKTFNPGAGLPTDKKKGG
	DVEAIKNAIANASTLAEV
	GLLQSGQIPGRERRLGPT
	EEEMEEDTVTNGS
	OGFPLLRRLKTLLVNNNRI
	EGLDQALPCLTELILTNNS
1 ! ! ! ! !	.GDLDPLASLKSLTYLSILR
NPVT	NKKHYRLYVIYKVPQVR
	FQKVKLKF\NPGAGLPTDK
KKGC	GPSPGDVEAIKNAIANAST
LAEV	/ERLKGLL
18957 49325 A 19066 92 303 NCFL	VFSPARSTVGEFASMSSE
	MHVVFLFPLTLCVVSLYER
	EKMFKGKRGAQLAKDIAR
RSKT	
[100.00 110	WWFPLGPTDDGEEEMEE\
DTVT	··-
10000	PWSTHASGWREATGFPQR
1 ; ; ; ; ; ; ; ; ; ; ; ; ; ; ; ; ; ; ;	AGCTMGKLTAE\LIEQAAQ
1 1 ! ! ! ! ! !	AVRDRELDLRGYKIPVIEN
	TLDQFDAIDFSDNEIRKLD LRRLKTLLVDTTRI\CGIGE
	A\LPCLTELILTNNSLVEL
	OPLASLKSVAY\LTYLRNP
1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	KHYRLYAIYKAPHVRVL
	VKLK\ERQKAEKMSKGK
, , , , , , , , , , , , , , , , , , ,	OLAKDIA\RRS\KTF*FPGA
1 1 1 1 1 1	*PKRKGGP/SLPGDVEAIQ
1 1 1 1 1 1 1	QNAFNSGLKVERLKGVCC
1 1 1 1 1	QIP\GRERRSG\PTDDGEEE
1 1 1 1 1	DTSHQTGS
	NEKIYELRVMETKP\DKA
	CDMNVDFDAP/LGYKEPE
1 1 1 1 1	QHEESTEGEADHSGYAGE/
	AFSGSGNRLDGKKKGVE
1 1 1 1 1 1	PIKPGDIKRGIPNYEFKLG\
EAGG	GRFVAFSGEGQSLRKKGR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
18961	49329	A	19070		1108	
18962	49330	A	19070	353	693	GSLFLVKRREPER/QVQHEELTE
10902	49330	^	17071		073	GEADHSGYAGELGFRAFSGSG
						NRLDGKKESPSPIKPGDIKRGIP
						NHEFKLGKTPFIRNACPLVKKF
						EEDEAGGRFVAFSGEGQSLSKK
18963	49331	A	19072	2	1112	RLGAAAAAAAGRAVGGVSSL
						HCPERSGVCQVVSIMFSFNMFD
						HPIPRVFQNRFSTQY\RCFSLS\M
						LAGPNDR/SSIGEKEGKIIMPPLG
						PGTKLSRL*HYRIPMLFK\LTNK
]		NSGPHGRHCGVAGSLWA**GA
						SAYLPHWDDGRTYSLGRKAA
						WVQGGRSVNLSSGPLISNFQPQ
						SP\DFL\DITNP\KAVFEN\ALRNF
		İ	İ			ALS*PPGDVI\AINYNEKILRNCV
						VMETQTPTRPVSHH*SVNH*TV
						EL*LLPPGATKEPRKDQVQP*G
						VRTEGESRPHSG\YAWKSLGFP
1						RFSRGSG\NR\LDGKKKGVEPSP\
						SPIKP\GDIKRGIP\NYEFKLGKIT
į						F/VSRNSRPLVKKVEEDEAGG\R
10064	10000	1.	10000	1,	416	FVAFSGEGQSLRKKGRKP
18964	49332	Α	19073	1	415	SGGGRNRSATGSWVGTMAGIT
						TIEAVKRKIQVLQQQADDAEER AERLQREVEGERRARE\RGMK
						VIENRALKDEEKMELQEIQLKE
						AKHIAEEADRKYEEVARKLVII
					}	EGDLERTEEPSLSWQESKCS\EL
						EEELKNVT
18965	49333	A	19074	1	421	
18966	49334	Α	19075	3	388	
18967	49335	A	19076	1	1332	
18968	49336	Α	19077	49	483	
18969	49337	Α	19078	227	1030	AVSWVGTMAGITTIEAVKRKIQ
						VLQQQADDAEERA\ERSQREFR
						EERRARE\QA\EAEVASLN\RRIQ
						\LVEEEL\DRAQERLGHCPCKKL
				1		*RS*KKLADESERRY*RLFEN\R
						ALKRLKEKIGTSREIPTSKKPKH
						HCRKEAD\RKYEEV\ARKLGDH
						LKEDLERTE\ERA\ELAESRC*EI
						DEQIRLIDQNLQCLSAAEDKYS
						QKEDKW\EEEMKILTDNLKEAE
						THAELAERSVAKLEKTIDDLED
						KLKCTKEEHLCTQRMLDQTLL
18970	49338	A	19079	50	864	DLNEM
18970	49338	A	19079	 	1014	
107/1	47337	IA.	13000	11	11014	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18972	49340	A	19081	1	1650	MGFHHVGQAALELLTSGFSQT
						QELQKFLFLLFLLVYVTTIVGN
						LLIMVTVTFDCRLHTPMYFLLR
						NLALIDLCYSTVTSPKMLVDFL
						HETKTISYQGCMAQIFFFHLLG
						GGTVFFLSVMAYDRYIAISQPL
				;		RYVTIMNTQLCVGLVVAAWV
						GGFVHSIVQLALILPLPFCGPNIL
						DNFYCDVPQVLRLACTDTSLLE
						FLMISNSGLLVIIWFLLLLISYTV
		1				ILVMLRSHSGKARRKAASTCTT
		1				HIIVVSMIFIPCIYIYTWPFTPFL
						MDKAVSISYTVMTPMLNPMIY
					-	TLRNQDMKAAMRRLGKCLKG
		1				PAVAVGPGPGPGDAEAAAEER
						RVKVSSLPYSVDALVSDKKPPK
						EASPVPAKSASSGATLRLLLLP
		ļ				GHGAREAHSPGPLIKPFETASV
		İ				KWENS\QDGAAWMQEPC*\YSP
		ł				PPRHK\HKTNPKPRTAFTTSQLL
		İ				ALEGKLLQKQYLSIAEGADFSS
						SPNLTETQVKILFQNRRAKTKR
						LQESELEKLKMAAKPMLPSSFS
						LPFPISSPLQAASIYAASYPFHRP
		ļ				VLPIPPVGLYATPVGYGMYHLS
18973	49341	В	19082	62	440	METERAL MECH CWAVA CA OVITA
18974	49342	A	19083	1	902	METEALMPGLGVAVAGAQVTL
		İ				GFLALETQLPFTGHKRILRLFFS SPRDPPHVPSKGLKRKWGARR
		ļ				GRTFALREAEPQRLPSLSRAFG
	* 5					TLARPYARLVRPGNPVVPFPLA
]						GSAEAARAAATCEROARTCPM
						PGVTVKDVNQQEFVRALAAFL
						KKS*V\GKLKVPEWVDT\VKLA
						KHKELAP\YDENWFY\TRAAST
						AQAPCTIRGGAG\VGSMTKIYG
			ŀ			GRQRN\GVMP\SHFS\RGSKS\VA
						RRVPPKPWRG*KMVEKGTKNC
				İ		GPQNLTPSRDKRDFGTEFAGQV
						AAANKEALEQIHCFGLIKLAHS
18975	49343	A	19084	1	673	AGPDAEASERVPEGPEAPGAGP
100,75	1,73,13	'				SRPLEPRPRASGRPGHGRCLGV
						TVK\DV\NQQEVRHELWAAFLS
						KSLR/LRVS*TVPEWL\DTVQAG
						QAQKSLLPYDE\NWFYHAELAS
						TA\RH\LYHPGSALGVGSMTKIY
						GGTSDETASMPKPLSARGFQEI
						WARRV\LQSPGRGLKMGGKRD
						QELRAAKLDTFKGQKRFWDRI
						RRNRLAVAYKEALEQTMAGV
						NKLPQFVKKKKK
			<u> </u>	L	<u></u>	1

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence		in USSN 09/540,217	location of first codon for peptide sequence		*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
		<u> </u>	1.000.5		Liaz	
18976	49344	Α	19085	l	427	HPVRFFVHRGPH/VDFSLEVVS
						QWYELVVFTASMEIYGSAVAD
	1					KLDNSRSILKRRYYRQHCTLEL
						GSYIKDLSVVHSDLSSIVILDNS
						PGAYRSHPDNAIPIKSWFSDPSD
				•		TALLNLLPMLDALRFTADVRSV
10077	102.45	1	10000	2	922	LSRNLHQHRL WNSAELGRGGPGAGGAGVIGM
18977	49345	Α	19086	2	832	MRTQCLLGLRTFVAFAAKLWS
						FFIYLLRRQIRTVIQYQTVRYDI
						LPLSPVSRNRLAQVKRKILVLD
						LDETLIHSHHDGVLRPTVRPGT
	İ					PPDFILKVVIDKHPVRFFVHKRP
						HVDFFL\EV\VSQWYELVVF\TA
						SM\EIYG\SAVGRLNWDNSRSIL
	Ì					*GGRY\YRQHCTLELG\SYIKDP
						LLWFHSDLS\GIVIL\DNSPGAYR
ŀ	İ					SHPGYGGR\DNAIPIKSWVSVTP
						ANTAILNLLPMLDALRFTADVR
		l				SVLSRNLHQHRLW
18978	49346	С	19087	69	206	
18979	49347	Α	19088	312	438	
18980	49348	Α	19089	635	826	QWISLWKCYKPEESGGQYLIFL
						KKSIFNDEFHVKPN*AS*GKEK*
			10000			TPLQTSNC*GILSPPGLP
18981	49349	C	19090	1	390	A D DI CCI A CA DED A DIAYANAYOT
18982	49350	Α	19091	336	551	ARPLGSLASAPFPAPM*NVYPT GP*\YNADCPNVTAPVCASNGH
						TFQNECFFCVEQREFHYRIKFE
						KYGKCD
18983	49351	A	19092	76	309	KGRFVIALTFTT*HSNLEKSCSK
10,00	17,551	1	17072		307	ICKIK\NKSLVCKHOLAKSIOSPS
						**WNGMNN*FTTLFICCHNTEK
						TFIHCGKRP
18984	49352	Α	19093	181	447	NKYQTDNIKLSDNDLTIGHQTG
						NTTAHC/RYIMLFRSFQQMAITT
						ILS*YRKNIYPLWKETLKRMWQ
						RHREKVGSFEKVNDIKILFHHW
1000	140252		10004		(20)	
18985	49353	В	19094	1	639	NI HEOMI WAIODID*I TEV*I DI
18986	49354	Α	19095	1	160	NLHSQMLWNQRIR*LTSV*LEL LCKNPNRVATRDQLLCLSGHL
						GMGRILELGG
18987	49355	A	19096	2	456	YKNTITDYYEHLYAHKLENLEE
10707	2,333	\frac{1}{2}	17070		1.50	MNKRRASTICTETIPKKLRRRD
						SSLTHSMRARRLSVDVKPQDRS
						RGPSECTIFRAGVPEGTRGALW
						LPHFQTESDPGSKPAITLLVIHP
						DELQIYVRTKTC/T/RMFIAALFI
						IAKT*KQPRYPSCLITST
	<u> </u>	<u> </u>	<u> </u>		<u> </u>	HAK I*KQPKYPSCLITST

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18988	49356	Α	19097	138	1193	LCFFGQHAPIYCSLRHSLLSYTR
						PASLICCLPGSHRYLEFVLLLLK
						HHRLHEDLNHCPIGSRSGRDVR
						SQTTQEQHVIAMPFPLRKLEPL
						HSGLYSHSCCSRYLAGWLALA
						TGGAARSPPAISILQPQEPSVTPS
						DTQRACGRSTEAFESKEQKASV
						RVETTVSFSCSIKVDGGLLEPKR
						ADTGWHGHHCRPASITPAGWV
				1		PARYLLYVFFSRQCVLSMVCGP
						QYTRGEIPGAVLKEECLPSLLT
						GRFDLSGRARRLSVDVKPQDRS
				1		RGPSECTIFRAGVPEGTRGVLW
						LPHFQTESDPGSKPAITLLVIHP
						DELQIYVRTKTC/T/RMFIAALFI
				· · · · · · · · · · · · · · · · · · ·		IAKT*KQPRYPSANTLLL
18989	49357	Α	19098	3	247	
18990	49358	Α	19099	1	1048	MQLKPMEINPEVSARCTATRRA
1						RGRGRGSRVADRFGFAFFAFA
1						FQMLNKVLS\RLGVAGQWRFV
						DV\LGLEEESLGSVPAPA\CALP
						ACLFPLTAQ\HENF\RKKQIEE\L
						KGQEVSPKVYFMKQTIGNSCG
						TIGLIHAVAQ*SKTNLGFE\DGS
				i		VL\KQF\LSETKKNVPLKDRAKC
						F\EKN\EAIQAA\HECRGTRKGQ
						CRVDDK\VNFHFYSG*PTVDGP
						PSMNLDG\RMPFPRGPMAPS\Q
						GDTLLEGRFPRSCRRISPRREQR
						RSPASSARGFSCKGSLNALWGG
		1				TWLISPLSPQHENILPPHASLKC
						FSTCETQAVLLFCQTRPSPQPHP
18001	40250		19100	57	450	RHLSTSRVHSCPLGHCGVSFRW VGDCLTSRGMSWVQATLLAR/
18991	49359	Α	19100	37	430	,
						ALCRAWGGT/CRGALTGTSISQ VPRLAP\RGLHCSA/ASVSSEQS
						LVPSPPEPRQR/PPGGERDKASF
						LQTVQKFADTSVVDSGHIDFIY
						1 7 7
						LALRKMREYGVERDLAVYNQL LNIFP
18002	40260	D	10101	207	200	LIMIT
18992	49360	В	19101	307	390	

SEQ ID	-	Met	SEQ ID NO:		Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
18993	49361	A	19102	431	1267	SGTRTTSTAARTSRTWRS*TPLL
		l	:			ACTPRPCGSAKLATATSRTCIRP
		İ				*SPSATQTLTVTKVPLPKDSTG
		ĺ				AADPPQPHIV/GAQGMYVSTPG
						IQSPDQQAALARHNPARPVFVE
						GPFSLWLRNKCVYYHILRADLL
						PPEEREVEETPEEWNLYYPMQL
		ĺ				DLEYVRSGWDNYEFDINEGKE
		ĺ				QHVTI/LCSSGEC*ILRRRQNV*R
		ĺ				LLDW/LAGKALTSPTCAPKITC
	•					VPLALPTPYCSGQNLNSSCSMT
	•					SVSDHLSPVPTLLSTISKAHGTP
						PPNTKVVTKVTVRQ
18994	49362	В	19103	49	691	
18995	49363	Α	19104	89	95	VGDCLTSRGMSWVQATLLARG
						LCRAWGGTCGAAL/SQEPPSLR
						SLASLPRGLHCSAAAHSSEQSL
]						VPSPPEPRQRAHQGSWCPLR/VP
						VLGQGAGLGERGQGRAFL/LD
						GCRNLRTTSVRKRGPH/VDLFY
						LALRKMRE\YGVERG/DLAVYN
						QLLNIFPKEVFRPRNIIQRIFVHY
		1				PRQQECGIAVLEQMENHGVMP
						NKETEFLLIQIFGRKSYPMLKLV
						RLKLWFPRFMNVNPFPVPRDLP
						QDPVELAMFGLRHMEPDLSAR
						VTIYQVPLPKDSTGASKIPPKPH
						IVGI/QRSPDQARPPLARHNPS\R
						PVF\VEGPFSLWLRNKVCV/YY
		1				HILR\ADLLPPEERVRA*SKACV
						GAGEWAGTTLRVHGRGWRRE
		1				TERALASDPACTPHLG\EEVEET
						PEEWNLYYPMQLDLEYVRSGW
						DNYEFDINEVEEGPVFAMCMA
						GAHDQ\ATMAKWIQG\LQETNP
						TL\AQIPRGLSAFAGSTRKLQTS
						SAGLEEPP/HCPRTTKEEDDNLQ
						RQSRAQS*SPCAGIRAQQQDRA
	<u> </u>					GLESAAGRVSR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
18996	49364	A	19105	l l	1199	EFGALRRTRLGSSFPRRRDSSA
						MESYDVIANQPVVIDNGSGVIK
						AGFAGDQIPKYCFPNYVGRPKH
						VRVMAGALEGDIFIGPKAEEHR
						GLLSIRYPMEHGIVKDWNDME
		l				RIWQYVYSKDQLQTFSEEHPVL
						LTEAPLNPRKNRERAAEVFFET
		1				FNVPALFISMQAVLSLYATGRT
						TGVVLDSGDGVTHAVPIYEGFA
						MPHSIMRIDIAGRDVSRFLRLYL
						RKEGYDFHSSSEFEIVKAIKERA
						CYL\SINPQKDETLETEKAQYYL
						PDG\STIETGPSRFRAPELLFRPD
						LIGEESEGIHEVLVFAIQKSDMD
						LRRTLFSNIVLSGGSTLFKGFGD
						RLLC*VKKLAPKDGRIRISAPQ
						ERLYSTWIGGSILASLDTFKKM
						WVSKKEYEEDGARSIHRKTF
18997	49365	Α	19106	1	3288	MPSGGDQSPPPPPPPAAAASD
						EEEEDDGEAEDAAPPAESPTPQI
						QQRFDELCSRLNMDEAARAEA
						WDSYRSMSESYTL/EGSGLWLP
						RGLFGHPGHCTGLLLLRSGCV
						RSQPSDRACR*EK\CHSSHLIEFF
		1				NKMKKWEDMANLPPHFRERTE
						RLERNFTVSAVIFKKYEPIFQDI
			•			FKYPQEEQPRQQRGRKQRRQP
						CTVSEIFHFCWVLFIYAKGNFP
						MISDDLVNSYHLLLCALDLVY
		<u> </u>				GNALQCSNRKELV
18998	49366	Α	19107	56	3552	GPAAQGCAMPSGGDQSPPPPPP
	,					PPAAAASDEEEEDDGEAEDAAP
						\PAESPTPQIQQRFDELCSRLNM
						DEAARAEAWDSYRSMSESYTL
						EGNDLHWLACALYVACRKSVP
						TVSKGTVEGNYVSLTRILKCSE
						QSLIEFFNKMKKWEDMANLPP
						HFRERTERLERNFTVSAVIFKK
						YEPIFQDIFKYPQEEQPRQQRGR
						KQRRQPCTVS*NFSIFCWVLFIC
	1					AKGNFPH*LSDDLVNSYHLLLG
	<u> </u>	<u></u>				ALDLVYGNALQCS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
18999	49367	A	19108	11	1193	GRRRFWWRKRIGGGGGMA
						MPAEAPWMASSFNGFLSLPPG
		l			•	WLLLPPPSDRNPPLQPSSRPPLC
		İ				VRMWMNAFTPLISLFFWASLV
						SSGARRLHPIQPQPCREQAPMP
						APGAARSAVAASMSDCTVAGP
		1				CTGSHTPCHSTSDSQSP/CGGVG
						SSLVV*AERSCAQCRMSPRKGH
						CCCCCCCCCC/SCCCYCHYC
						ccyccccccccccccc
						CCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC
						CCCYCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC
						CC*HYCCCCCCF/GCCCCRYC
						CCCRYRRCYCFCCCCCCCCYCY
		1				CCC*RTAHAWPSMNKSQCVLP/
						CICKVPLGRLAVTTGCPGFG*G
			1			GCWEDCMFPAKGPSIRLEQWFS
		1				TTAILLPSGPGDFWQYLGTLLV
		1		ļ		VTTREEVLLESGG
19000	49368	Α	19109	1	762	GTRDATAEENRVLLAMVNPTV
						FFDIAVDGEPLGRVSFEVRGLD
				1		TKK*LLI*SIKLC*QIGGSSIFITS
		ļ				D*KNSCLPLIVQQCLLFLRILP\L
		ļ				FADRVPKASQKIFVALS\TGEKG
		ŀ				FWL*GVPCFHRLFP\GFYVSRGG
		1				DFHTAINGTG\GKSILWGENLK
		ŀ				DENFI\LKHTG\PGI\LSMANAGP\
		ļ				NTNGSPVFLTSCTA\KT\EWLDG
		ŀ				K\HVVFG\KVKEGMNIVEAMER
		ŀ				SLGPRMGKT\SNKFTIADCGQL
:						RIKFDLVFYL
19001	49369	Α	19110	95	240	LHSEITVCL*GSEHHFFENGTYI
		<u> </u>				PHRSCHDPLEARAEDFL
19002	49370	Α	19111	1	1644	
19003	49371	Α	19112	237	632	AIGFPSYSSMQLVNIFGLLNLFSI
						ISSSFRHQG*HS*ELP\YPLEPGA
						EDFLR*GP\LAGHGHTGSQPFFC
		}				RVSEA/CGSHLVNSSRAALKRW
						AGV*RKIWPRGSPETPRNLKDL
						SVLEYENTPDTSGKKALED

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
19004	49372	Α	19113	274	2704	GISVLHGCGYRFQPRGMAAPGP
		1				FSEKPVPGCDAGDLQTPALSSW
		1				TQTFIISHEQGIKFLNQRWSCWS
						KERNHGHCRVRGHVTAAQCDL
						FGNGLKHNLDLHIHDRNDASK
						NVKKSTNYGKMSFYTYCECTP
						TGEKLWDHNQHRKIIGYKPASS
			!			QDQKIYSGEKSYECAEFGKSFT
						WKSQFKVHLKVPTGEKLYVCI
						ECGRAFVQKPEFITHQKTHMRE
		ŀ				KPYKCNECGKSFFQVSSLFRHH
						RIHTGEKLYECSECGKGFPYNS
						DLSIHEKIHTGERHHECTDCGK
						AFTQKSTLKIHQKIHTGERSYIC
:		•				IECGQAFIQKTQLIAHRRIHSGE
						KPYECNNCGKSFISKSQLQVHQ
İ						RVHTRVKPYICTEYGKVFSNNS
						NLITHEKIQSREKSSICTECGKA
						FTYRSELIIHQRIHTGEKPYECS
						DCGRAFTQKSALTVHQRIHTGE
						KSYICMKCGLAFIRKAHLITHQI
						IHTGEKPYKCGHCGKLFTSKSQ
						LHVHKRIHTG\EKPYMCNKCGK
		ŀ				AFTNRSNLITHQKTHTG\EKI\FI
						CSKCGKAFTQRSDLITHQRIHT
						GEKPYECNTCGKAFTQKSNLNI
				1		HQKIHTGERQYECHECGKAFN
				i		QKSILIVHQKIHTGEKPYVCTEC
						GRAFIRKSNFITHQRIHTGEKPY
ļ						ECSDCGKSFTSKSQMT\EH*A\V
						HSG*KPYV\CAECGKAFSGRSN
						LSKHQKTHTGEKPYICSECGKT
19005	49373	Α	19114	3	180	
19006	49374	Α	19115	95	433	PVLRTHPGPQSLPRVPGVPCGG
						LLEPLSRAEVSPRLGLRRDLLG
						GMAPSGSST\VF\LLALTIIASTW
						ALTPTHYLTKHDVERLKASLDR
				1		PFTNLESAFYSIVGLSSLGAQVP
19007	49375	Α	19116	196	706	ISISNETKOLLLAAVSEDSSVTQI
						YHAVAALSGFGLPLASQEALSA
						LTARLSKEETVLATVQALQTAS
		1				HLSQQADLRSIVEEIEDLVARL\
						DELGGVYLQF\EEGL\ETTALFV\
						AATYKL\MDHVGTEPSIKEDQV
						IQLMN\AIFSKKNFE\SLSEAFSV\
				1		ASAAAVLSHNRYH

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19008	49376	Α	19117	275	2192	MVVSPQGSSTVFLLALTIIASTW
-						ALTPTHYLTKHOVERLKASLOR
						PFTKRLCFSWFTTCVTT*H*LFP
						TV*KACTYIRSKSLPCLPQIAKT
						CSKTLCLPILS*ISISNETKDLLL
]				AAVSEDSSVTQIYHAVAALSGF
						GQVREGVSG*NKTCHSLGFYSL
						PSRTVDSGSCISFLLLLDIQHISV
						LLWQDLVARLDELGGVYLQFE
						EGLETSTAVALPPSQQRKRNSCI
						SHFFQDQVIQLMNAIFSKKNFE
						SLSEAFSVASAAAVLSHNRHPR
						NKILCSCSLCCNK*FFIYFQLQV
						TNVLSQPLTQATV\KLLFVISMT
						F/RT*VLAKPIVILNRDVFELNF
						MNVKFYICM*IPEHPPIDLATVF
}						SIFL*LRVKISTDIFSACSQRYTD
						PL*RLTLYASFRVTYPAKAKGT
						FIADRDSMLGPGRILKLSDERL
						YYSSI*TFVRLHNQKTGQEVVF
						VAEPDNKNVYKFELDTSERKIE
						FDSANIIY*VSPTSLPLCFYRLSL
						FF*ADVVINPGIHGCPYWEGHIL
ŀ						KSSWCVFRQHLFREPEKSWHG
						LCCFPFKF*CLFNLFLPIIQWILT
						FLNRRASLCFI*LDVMTPLLAA
1						MLGLMYVYWTQLNMFQTLKS
ĺ						LAILG\SVTFLA\GN\RMLA\QQA
19009	49377	Α	19118	3	413	PRKEAPSTFLSQ/NLFTPKQEIQ\
						HLF\REPEKR\PPTVV\SNTFTALI
						LSP\LLLLFALWIRIGA\NVSNFT
						FGLPSTI\IFHWG\HAAMLG/LSM
		ŀ				YVY*TQAQPCSQTLEVPWPILG
						QCDRFLAGQSGMLAQRTAVQE
	İ					NQHN
19010	49378	Α	19119	2	535	GRVDPKAKKEAPAPP*SLKPKA
						*AFKGPRKAVLK\GV\HSPQKA
1						GRSRTSTPPFRAGRKTTADSRR
						PAPKIILRKSRVPGGNKLD\HYA
						\IIKFPADH*SASAVKKIEDN\NT
						L\VFIVD/VLKANKH\QIKQGCE
						RSLYDI\DVGQGQHPWIRPD\GE
						KKAYCSTWLLDYDGFGMFAN
l						QNWGSI
						GRSRTSTPPFRAGRKTTADS PAPKIILRKSRVPGGNKLD\H \IIKFPADH*SASAVKKIEDN L\VFIVD/VLKANKH\QIKQG RSLYDI\DVGQGQHPWIRPD KKAYCSTWLLDYDGFGMF

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	İ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19011	49379	Α	19120	159	801	GAAGRRRLPEGEGEDEASAWR
		-				RLARCLMPKKNRIAIYE\LLFKE
						G\VMVAKKDVHMP\KHPE\LAE
						VKNVPQPFMVMKGHAGLSSPR
		İ				G\YVKGTSFA\WRHFLTGTLTQ*
			}			GVSQYSPWITLHLAPGRFVPCP
						PLRPVARPRRMGRP\RPKKVLG
		ŀ				GVEATWRGFHKRGEV\DRGYL
		ļ			!	TRRSACCQLGA\DKKSRGWGL
		ŀ				GSSNPNFQFRG\GF\GCGRGQ\PP
19012	49380	Α	19121	334	989	EPPAETLPVAARDPTGLSPTYLS
						SHKHEKFQQVRHFPVLLGYPIA
		Ì				SSGASPEVRQRRTTFFRFRPGES
						LCGDMKLLTHNLLSSHVPAGG
			İ			SLPFSGGMVGRGPVLA*TP/LAL
						QATEGR\ICPVEINPQLRGRVMI
				1		P*KWKWS\AF\LEAGPDKLAV*I
						PGARKGPG*GDMRENEEFS*GP
						WHHLACWEVGSD*EGTL\QVP
						GNLGRYFSPFKPRGSPNMLAE
19013	49381	C	19122	61	177	
19014	49382	Α	19123	4	191	
19015	49383	A	19124	1902	2187	
19016	49384	Α	19125	55	340	GELVLKALSAFIRAFRCPAEGS
				1		DPVGRAADP\SPPAPRGISQGGS
						ACGSPFLPGPITESLPLPPGLGSG
						SELPSLYLVSGIYGGVRRGPPA
	10000	ļ	10101	1000		RLEA
19017	49385	A	19126	1228	2447	TSQGSSPPAHIGRWSSNHKKGL
						GSSFSQSCCKAEALVKNAWPPT
						QA*VSICCFEQDLGRGQQKGKS
						GGKWQKTAESREAVTECPACY
						RAARCRAAGDRGLRAEGAAAG
					<u> </u>	HTASPSAYRVRGERGHRDMHS
						ASARLSHPGETNLMLGLAEPSV
						TSH*SHPFRQS*LP*QAGP**EM/
						PLPPSARDSSDLCSG\SPSHDLLP
		ŀ				LPNTVIFWGPSSPQDWQPLPLSP
						FCVSQAAGSPCSPLLQGLEPGGI
						STSCPPGPAPYIDFGAVSGAGQS
						AAQRGAHFPAAMAERPSAGTS
						GSSGISGAIAATVVSWVAEGEQ
						AVKCLAEGSTRQTGQPSRRAPS
						RGCPAPGAPPGRAAPPQAPPAA
						PAAGGGRGPPAAPRAPAPRPQP
						QGAPRPVSPCRGTGAAPPPTLV
L	<u> </u>	<u></u>	L	<u> </u>		WGLSSEAEG

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	1	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19018	49386	A	19127	164	563	CEAALKGDSGGSPWPDEKPKE GVKTENNDHINLKVDGAEMVT VV\QFKIK\RHT\PL**T*LKAYL* TDRGLSMRQIRFRI*PGNPIN*K QTHLHQLGNGRIEDYNLMCFQ QARPGGVYLKKGTCFFYSKNS VSF
19019	49387	В	19128	166	426	
19020	49388	С	19129	76	285	
19021	49389	Α	19130	3	179	
19022	49390	A	19131	93	172	
19023	49391	Α	19132	1	2742	
19024	49392	Α	19133	1287	2571	ILRQWLLASPEEKTGRPALASG SAETLSWLRAQGREPRGPA*SA VGAEHRTRTAPGSDPRCGILPS GRSPPSP/PARLFLASVPATI*AM TPSPTRLPAWPPPCRG/PDASPE VVRKGGKAGQPRGSPQPWRSG AEEIVEVGLLPLTPRALLFPQVP TALPAVYHFLAACEGICPHSCA APSMIELWDFLIQQHQPEKAVA CTTTHIPGLRISALNKHRAWGG TDDDVAASEMPLGAAGAEPW D/VE*LLPAALKIPTGHTATTSW GAGP*SPQGSCKG\SGVGSGAT ADLLAPTAGVGTSRTACGGGT QFMSEGAPGVMNNGSPDWSST QFMSEGAPGVMNNGSPDWSSN LSQPQPMHLLYLTTWFLERPM CCPDTTALTQTLPVPSFSYPVCT SHIRQNPHLHFLRAPSPHYTTA CSFEPQRFLEQQQDLL
19025	49393	Α	19134	31	202	ILITYIIKLINQADFSTP*FMSHLI VSSRILCTEKWNFHFPPHF*D*Q QTFCTLQFL
19026	49394	Α	19135	410	867	LLTLSRTTPLMSLGFPGIPKMIN RPRQSSPIKLQNSSSLSNLLFFQ EDSSTLDSGLERSQGLDSTGGG EDICRVWTAS*SFICLSDPQKVQ IQSAPNRKAWCLVILVVVSHSS TPFSWQKVMKWIKKISFRVFAA GQGLLGGYSNPRYSFSF

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	1	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	i	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19027	49395	Α	19136	111	743	VITILTPMLADRTRRIERPPKKK
	1					GTTSLGQRKWLLTQDWPSVYP
						VA\HPFKPSAVP\LPV\RMGYPV
		ŀ				KKGRAHGDRRGNLELLKISQFF
						WILTSCSNLKSTCEALKDF\CTG
						VGQPHWDSGRKKMLKKHFSN
1						WKFGQHLDLCFHQGPSGSGNP
						RARVVVLKSKAFLV*I*HDHAK
						KTLITLVGERYCKTPDVLTIKQ
						NRWPLRSQITIMPVYL
19028	49396	A	19137	305	2291	YLDAEKMGQKASQQLALKDSK
.,,,,						EVPVVCEVVSEAIVHAAQKLKE
						YLGFEYPPSKLCPAANTLNEIFL
						IHFITFCQEKGVDEWLTTTKMT
						KHQAFLFGADWIWTFWGSNKQ
						IKLQLAVQTLQMSSPPPVESKP
						CDLSNPESRVEESSWKKSRFDK
						LEEFCNLIGEDCLGLFIIFGMPG
Į.						KPKDIRGVVLDSVKSQMVRSH
						LPGGKAVAQFVLETEDCVFIKE
		1				LLRNCLSKKDGLREGGASPGSL
						RLAAPGPPLTLNAACPLRLAVL
						AAMAAAALPAWLSLQSRARTL
						RAFSTAVYSATPVPTPSLRVDD
						LHLTEIVGMLDSVLTPEDSSGK
						YRFISGEVLCRITGCFTGVRVEA
				1		KDLFGGCCSNPNEVMVTWIKVI
		1		1		VEKEVWLYLRYILKALPPRTEK
				1		MAVDQDWPSVYPVAAPFKPSA
		1				VPLPVRMGYPVK*GVPMAKEG
				1		NLELLKIPNFLHLTPVAIKKHCE
						ALKDFCTEWPAALDSDEKCEK
		1				HFPIEIDSTDYVSSGPSVRNPRA
						RVVVLRVKLSSLNLDDHAKKK
						LIKLVGERYCKTTDVLTIKTDR
		1				CPLRRQNYDYAVYLLTVLYHE
	İ					SWE\TEEWGKK*D*SRHGKSIY
						GENSSSERKYPGERFSR*KLLR
		1				KNMGN*LKKSS/CGTKEIEEYK
						KSVVSLKNEEENENSISQYKES
19029	49397	A	19138	3	233	KS V VOEK VEEE VEIVE () KES
19030	49398	A	19139	1	256	
19031	49399	С	19140	5	114	
19032	49400	Α	19141	8	444	TTDKEQ*EL*LLLDTE*GSYTKE
						CHTDSLSRY*TSSTSAEDRDDSL
						LRRSGSYRDL*E*IPYSCRLDKD
						DSTDCIKLYEQILAE\NDKLKAQ
						LHDTNMELTDLKLQLEKATQR
				1		QERFADRSLLEMEKRERTALQR
				1		RISEMEERLQM
19033	49401	A	19142	1	287	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19034	49402	Α	19143	211	346	
19035	49403	Α	19144	175	3381	EDPAAGEGMKMADAKHKRNE
						QLKRWIGSETDLEPPVVKRQKT
						KVKFDDGAVFLAACSSGDTDE
	1					VLKLLHRGADINYANVDG/LSH
						ALHQACIDDNVDMAKLLVDNG
						ANINQPVNDGWIPLHAAPSCGY
						LDIAEFLNGQGAHVGAVNSEG
	İ					DTPLDIAEEEAMEELLQNEVNR
						QGVDIEAARKEEERIMLRDARQ
						WLNSGHINDVRHAKSGGTALH
						VAAAKGYTEVLKLLIQAG\YDV
	}					NIKDYDGWTPLHA\AAH\W
19036	49404	Α	19145	1	2406	
19037	49405	Α	19146	1154	1365	
19038	49406	Α	19147	1	293	MGVSVLDFSHSDKCEVASRFAS
						GMPEPTNIVLPKLEVADTLGGC
						TTIPESDLEERSVEQDSTELFTN
İ						HRHLTAETPR/PW*RLDGLHKQ
		ļ				VSPLQGVSE
19039	49407	Α	19148	3	1290	VSQATDVEVGTDLVPSVTVKV
						TLQNRVILQKAKLSVYVQPPLE
						LTCDQFTFEFMNRNPDGIPRVIQ
						CKFRLPLKLICLPGQPSKTASHK
						ITIDTNKSPVSLLSLFPGFASQSD
	1	İ	;			DDQVNVMGFHFLGGARITVLA
	į					SKTSQRYRIQSEQFEDLWLITNE
ŀ						LILRLQEYFEKQGVKDFACSFS
						GSIPLQEYFELIDHHFELRINGE
						KLEELLSERAVQFRAIQRRLLA
						RFKDKTPAPLQHLDTLLDGTYK
						QVIALADAVEENQGNLFQSFTR
ŀ	ŀ					LKSATHLVILLIALWQKLSADQ VAILEAAFLPLQEDTQELGWEE
						, , ,
						TVDAAISHLLKTCLSKSSKEQA
					}	LNLNSQLNIPKDTSQLKKHITLL
						CDRLSKGGRLCLSTDAAAPQT MVMPGGCTTIP\ESDL*ERSVEQ
						DSTELFTNHRHLTVETPRPEVSP
19040	49408	A	19149	48	642	WNSALREPRLNGADMAKSKN
19040	49408	^	19149	140	042	HTTNNQ\SRKWP\RNGIKKP\RSP
						KIRILLKGVAPPSFL\RNM\RFAQ
						ESPNKKGP*RRLQANKLPRAMS
						ATLPRAIKA\LVKPQGGLSPKIP\
					:	KGVSSRKLDSTCLTFAQPQALG
						SVARCP*LPRGPRL\CPPKAKAK
						AKAKPRPKPRPRAKDQNQGPG
		1				LQPQASVPAQ\APKRTQAP\TKA
L	<u> </u>	1	L	L		SE

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19041	49409	A	19150	1953	2447	WTPQYMDTEYKNGPWMMEK KVWLDP/NETNEIPNANFRQQIR KLIKDGLILRHRKPVTVHSRAQ CWKSTLARRKGRHLGIESKKID RHMYHSLYLKLKGNVFKHKRI LTEHSHKLKADKARKKPLADQ AEARGSKTKEARKLREEHLQT
19042	49410	A	19151		693	KKEEIIKTLSQEEKAKK MAVMEKLYWVNREESYASLR VHLPICNATTMSMLSLQKRLAS SVLRCGKKKVWLDPKETSEIAN ANSHQWIRKLIKDGLIIHKPVTV HS/*ARYR/KNTLACQKGRHMG I*KATANARMPEKIMWMR/RM RILCQLLRRYCDSKKIDHHMYH RRYHESKKVDRHMYHSLYLKV KGNVFKNKRILMEHIHNLKAD KACKKLLADQAETRRSKIKEAR KHREEHLQAKKEEIIKTLV
19043	49411	A	19152	34	280	EVIHAAIGEKKGSY/NA**PWVE EQLTRQPLIHHQPASLHSVY/RC RYHSLYLKVKGNVFKNKRILM EHIHKLKADKARKKLLA
19044	49412	A	19153	2	625	HEAMSMLRLQKRLASSVLRCG KKKVWLDPNETNEIANANSRQ QIRKLIKDGLIIRKP\VTVHSRLR CRKNT\LAR*KGRHMGIGKRK VPANA\RMP\EKVT\WMRENEG FCRRLASEDTRES\KKIDRPHVV TALYL\EVKGNVFKNKRIL\ME HIHKLKA\DKARKKL\LAD\QAE ARRSKTKEATKRREERLP/ARPR KREII*TLSKEEETKK
19045	49413	С	19154	154	291	
19046	49414	Α	19155	3	253	
19047	49415	A	19156	2	474	FKDKQNKPTGFALGSIEGRVAI HYINPPNP*V*LCQLADFTGLFL TKEDLHEPLSFSAKDNFTFKCH RSNGTNTSAPQDIYAVNGIAFH PVHGTLATVGSDGRFSFWDKD ARTKLKTSEQLDQPISACCFNH NGNIF/AYASSYDWSKGHEFYN PQKKK

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19048	49416	Α	19157	97	1728	IRGFFRGSQGSFYRRLKEASGS
		ĺ				WDFCPRSWPSSLRARSGSQTPE
						TLLLPAGVFRRSETPMFKMSLF
		ĺ				GTTSRFGTSGTIMFGSAT\TDNH
						NPIEGY*K*HSFSLMDSIG\CSCL
						FSPPTLPGNFLIAGSW\ANDVRC
						W\EVQDSGQTIPKAQQMHTGP
						VL\DVCWE/SDGGSKVFT\ASCG
						\KTAKMW\DLRQ*PRRLQIAQH\
						DAPV\KTIHWIKAPNYSCVMTG
						SWDKTLKFWDTRSSNPMMVL
						QLP\ERCYCADVIYPMAVVATA
						\ERGLIVLIQL/ERIQPSEFRRIESP
						L\KHQH\RCVAIF\KDKQNKPTG
						F\ALESIQGRVAIHYINPPNPA\K
						DTFTFK\CHRSNGTNTSAPQ\DI
						YAVNGI\AF\HPVHGHPCNCGD
						LDGRFSF\WDI\DARTKLKNFRE
						QLDQP/LSSAC\CFNH\NGKHILP
						YAS\SYDWVKGDH\EFYNP\QK
						KKFTFFPA*MQAEELKAPGNKE
						VVAGRTLGSAQSLFLLHSGLIS
						VRIWAPNLVGVVSPWTMEFQP
						PGENDVIVQQLESPRRPAGDLP
						VFSIPTGLVAEVFSGKPKGGLK
19049	49417	Α	19158	1	1254	
19050	49418	Α	19159	1	921	
19051	49419	Α	19160	1	1386	
19052	49420	Α	19161	49	731	

2381

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19053	49421	Α	19162	61	1533	PLKAKMGKEKTHI\NIVVIGHV
Ì						DSGKSTTTGHLIYKCGGIDKRTI
						EKFEKEAAEMGKDSFK\FAWV
						LDKLKAERERGITIDISLWKFET
						SKYYVTIIDAPGHRDFIKNMITG
						\TSQADCAVL\IVAAGVGEFEAG
					:	IS\KNGQ\TREHALLAYTLG\VE
						QLIVGVNKM\DSTEPPYSQKRY
						EEMWLREVSTF\IKKIWLQPPTQ
						*HFVPISWFGIGDNM\LEPSA\N
					:	MPWFKGW\KVTRKDGNASGTT
ļ						LLEALDCILPPTRPTDKPLR\LPL
						QGVH\KLGGIGTVSSAPMET\GF
						SNPGMVVTFAP/SSTVTTESKIL
						SKMHP*SFEVKLLPGDQCGPSM
	1					SKNVFCQGMFRRGKRLLGDQQ
	İ	1				KMTPPNGKAAGLHWSRVIIL\N
		1				HPRPK*GAGPMPLYLDC\HTGS
	ļ					HLHAKFAELKEK\IDRR\SGKKL
						EDGLKFLKSGDAAIVDIVPG\RP
		ŀ				MCVESFSDYP\PLGRFAVRDMR
	}					QTVAVGV/ISKAVDKKGCWEL
						GKVTKVWPRKAPEG
19054	49422	Α	19163	1	514	EFGTRAKKEAPAPP*SLKPKRK
						AFKGPRKAV\LKGV\HSHQKEE
						DPARSPTFPGGPKTLR\LRRQP\K
						YPRKSRVPGETRLDHLCLSI\NF
						PLTH*VLPMKKIEDNNTLVFIV
						DVKANKH\QIKQAVKKL\YDI\D
						VAKVNT/LWIRPDGEK\KAYVR
					İ	L\APDLPIAFGMFANKIWGFI
19055	49423	Α	19164	433	1015	GLISNLDMASRGAS*\PTGPDTG
						NKICQFKLSTSGESPAVCKSSLV
						LRFVKGQF\HEFQESTIGAAFLT
	-					QTV\CLDDTTEKFEIW\DTAGQE
						RYHSLAPMYYRGAQA\AIVVY
						DITNEESFARAKNWVKELQRQ
						ASPNIVIALSGNKADLANKRAV
						DFQ\EAQSYADDNSLLFMETSA
						NTSMNVNEIFMAIGKINISF
19056	49424	В	19165	1	2511	
19057	49425	Α	19166	2	360	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19058	49426	A	19167	125	211	FRPGSPRQPRAQPISAPDCTRA MVGRRAL\IVL\AHSERTSFNYA MKEAAAAA\LKKKGWEVVE\S DLYAMNF\NPIIS\RKDIT\GKLE GPLRTFQYPGRSLFLAYK\EGHL EP/EIICGLNKKKLEAADLV\IFQ FPLAVGLEVPAIL\KGWFEPSVH KESFA*HFTAGHVMDKGPFPG VKKAVAFPFTTGG\SGS\MYSC QGI\HG\DMNVILWP\IQSG\ILPF LGLPKSLDPQLT\YSIWPPLPAR RPEFSILE\GWKK\RLE\NIWDET\ PLYFAP\SSLFDLNFQAGF\LMK KEVQDEEENKKF\GLSVG\H\HL GKSIPTD\NQIQS*KRECTVHYT RDSHKVAAGAAQLTESLVPAR VAPATTSPANQRPGLHQSHGR
19059	49427	A	19168	2	331	WVDSQQK\RYVPVKGDHVIGIV TAKSGDIFKVDVGGSEPASLSY LSFEGATKRNRPNVQVGDLIYG QFVVANKDMEPEMVCIDSCGR ANGMGVIGQDGLLFKVTLGLIR
19060	49428	A	19169	145	350	QEFHLSSEPSIRLGRSLYHIHHL FASKKVLD*QWLDQSL*VCSW EHMRRSQ*GIFSRLANRGFQNG DC

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19061	49429	A	19170		1863	MAVAIRLSADTQWEDSPGKSIL ELHMSCTKIPERCEADKKESTS SWCTRKIRFRMDLEKECKVLLS GGGGSRMGSQKGDGVGRWSSP RVGLPSSQTLLRPPPTDFHIAPL SKARQHLLVSVSVLFCFSAPLD VQPLKVLDECQNQRACHLLVN SRVFGPDLCPGSSKYLLVSFKC QPKQMFTQQGPGPVLSIGETQV STSSVTRSSSPVGPDIHCEFHTN MCKIPDELKNKTVCEDQELKL HCHESKFLNIYSATYGRRTQER DICSSKAERLPPFVFTRYYCFLY APVSSIANPVKGCFHLGKKQDC LSYSALQVLSRRCYGKQRCKII VNNHHFGSPCLPGVKKYLTVT YACATLSSMGPILAEVPPEAKK VTLSTSLDSSSGRMEDGRLDTG HTEVPVTSGLFIRVLNKKQNFS LQIRCSFGRGPGKLRHCWAGA AIALGTEYILVKGDQVTGIVTA KSGDIFRVDVGGSEPVSL\SYLS FEGATKRNRPNVQVGDLIYGQF
						VVANKDMEPEMVCIDSCGRAN GMG\VIGQDGLLFKVTLGLN*E SLLASQICE/ISIQEVGKL\LPLEI VFGMNGRIWG*GQKTHPSQTLI
						LA\NIL\EACEHMT\SDQRKTDLP
19062	49430	В	19171	29	82	
19063	49431	Α	19172	1	427	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19064	49432	A	19173		1548	MTMAAAAVVARGAGARAATA AALRGGCGTAARGRPCAGPAR PLCTAPGTAPDMKRYLWERYR EAKRSTEGTKKYTTSLNARHY YTHFTEENEDEINSSSSYASQKK TFEINPRHPLIRDMLRRIKEDED DKTVLDLAVVLFETATLRSGYL LPDTKAYGDRIERMLRLSLNID PDAKVEEPEEPEETAEDTTE DTEPDEDEEMDVGTDEEETG KESDDPMAYIHFTAEGEVTFKS ILFVPTSAPRGLFDEYGSKKSDY IKLYVRRVFITDDFHDMMPKYL NFVKGVVIRKKLVRKTLDMIK KIADDKYNDTFWKEFGTNIKLG VIEDHSNRTRLAKLSRFQSSHH PTDITSLDQYVERMKEKQDKIY FMAGSSRKEAESSPFVERLLKK GYEVIYLTEPVDEYCIQALPEFD GKRFQNVAKEGVKFDESEKTK ESREAVEKEFEPLLNW\MKDKA LKDKI*K/ALWVSSAALTESPVL LLVAQPVRDWS\GQHWERIMK
19065	49433	A	19174	37	266	AQAYQTGKDISTN\YYA WADRAAGGVRILTQGCVGGG GSCDRRGLETSPARTPVRALWV LG\LCCVLLTFGSV/READDEVD VDGT\VEEDLG*KYEKDQVTDD EVVQREEEAI\QLDGLNASQIRE L\REKSENFPF\QAEVNRMIETLF IQFHCIKNKEIFLERLISNSSC\AL DRLSLISLT\DENALSGNEELTV K\IKCDKEKNLLHVQHTGVGM T\REELVKN\LGTIA\KSGTKRVF *TKRPEA\QEDGPSQPS*IDWPS LGVRF
19066	49434	A	19175	233	664	RAPEQSSGVSAAEQERVPNAG WPCSVAPTCPSPRSAEPARCT* GKEEGG/CEDSDRADSGSPFQE PRYPSSGPFPIPQASFPCPCLLVL GIRGHSTILQYRGQEHHVLFTG RCLLVDPTLLVLVVSKKSPNAA RDQPARYPRGPL

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19067	49435	A	19176		672	MAEAGPKPQRTLMWAWESVL SQRKPREAGRLSEGCVKATGTP GFGWQARPDQKEHRLDGRVID PKDRHGLVKDTEPRLNEQSLCG GLNPEATEEKIREYLASLGRRR PVKKVLEKKFHTVSGS/KCGPQ VPPSWSETYRTRA*SGDQLGTP AAPNPRWPME*EQEIEAQRRSA HSGARAPWPPWPPRPDGPPGNP GPKFMVSLSCWNISSLRAGLLY AGCGCSP
19068	49436	Α	19177	3	684	RHASARKKYNPPSWFHRGLPPP EDQEDEMSVSEELE\PITLTPSSA LKPSDRMTMSSLGERACCRDY QRLGLGTLSSSLSRAKSEPFRIS PVNRMYAICRSYPGLLIVPQSV QDNALQRVSRCYRQNRFPVVC WRSGRSKAVLLRSGGLHGKGV VGLFKAQNAPSPGQSQADSSSL EQEKYLQAVVSSMPRYADASG RNTLSGFSSAHMGSHGKWGSV RTSGRSSGLG
19069	49437	Α	19178	1	341	
19070	49438	A	19179		3710	MVSGARAAAAAAAAAAAAAER TRGVPGHPDSTGRLPAHQQLQ RGLLFRQLGITNVLSLVCATLM EHEVLFLSRSYQQLTDACRSLL ALLFPLRYSFTYVPILPAQLLEA LSTPTPFIIGVNAAFQAETQELL DVIVADLDGETVTIPECVHIPPL LEPLQSQTHSVLRMVLDPELEL VDLAFPPPM/HDKELRLFHQLL QGYLWCLHV\VCIHPEPVIRFH KAAFLGQCGLVEDNFLMNVLE GMAFALFVLECGVP
19071	49439	A	19180	1	303	TVI TCCDC*I A*NDCM/HNNCE
19072	49440	A	19181	14	942	TKLTGSRG*LA*NPCM/HNNCE SCVDLLFVRGAGNCPECGTPLR KSNFRVQLFEDPTVDKEVEIRK KVLKIYNKREEDFPSLREYNDF LEEVEEIVFNLTNNVDLDNTKK KMEIYQKENKDVIQKNKLKLT REQEELEEALEVERQENEQRRL FIQKEEQLQQILKRKNKQAFLD ELESSDLPVALLLAQHKDRSTQ LEMQLEKPKPVKPVTFSTGIKM SVGITLCPAEEHTFNDERSFNLS SSWSKFNKPPSTLPFKVLDEER LVELKVPNTLITKLVPFAISPCS EAIQDHPAILLAYKKTPFWRLQ
19073	49441	В	19182	49	456	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19074	49442	В	19183	53	333	
19075	49443	A	19184	2	208	VYPELQITNVVEANQPVTIQNW
						CKRGRKQCKTIPHL*FPYRCLG
						EPAGRGAGVDWGPGLEGRKRG
						csc
19076	49444	В	19185	51	300	
19077	49445	В	19186	148	3568	
19078	49446	A	19187	1	3411	<u>.</u>
19079	49447	Α	19188	2	2416	NSRGAACAGPRETAAVAARAE
						QGRGGSHSHSSALGAPRRVAM
						LPGLALLLLAAWTARALEVPT
						DGNAGLLAEPQIAMFCGRLNM
ŀ						HMNVQNGKWDSDPSGTKTCID
						TKEGILQYCQEVYPELQITNVV
						EANQPVTIQNWCKRGRKQCKT
l						HPHFVIP\YRCLVGEFVSDALLV
•						PDKCKFLHQERMDVCETHLHW
						HTVAKETCSEKSTNLHDYGML
						LPCGIDKFRGVEFVCCPLAEES
						DNVDSADAEE\DDSDVWW\AG
						ADTDYADGSEDKVVEVAEEEE
					İ	VAEVEEEEAD*LTRTDEDG\DE
						VEEEA\EEPYQEATERTT\SIATT
						TTTTTESVEEVVREVCSEQAET
						GAVPSNDLPLVL*CD*REVCPIL
						LRRMWRQPEQL*HRRVLHGRV
						WQRI\PTTAASTPDA\VDKYLET
						PGDENEHAHFQKAKERLEAKH
						RERMSQV\MREWEEAE\RQAK
						NLPKADKKAVIQHFQ\EKVESL
						*QEAANE\RQQLVETHMARVE
						AMLNDRRRLALENYITALQAV
		1				PPRPRHVFNMLKKYVRAEQKD
		1				RQHTLKHFEHVRMVDPKKAAQ
		1				IRSQVMTHLRVIYERMNQSLSL
						LYNVPAVAEEIQDEVDELLQKE
		1				ONYSDDVLANMISEPRISYGND
[ALMPSLTETKTTVELLPVNGEF
		1				SLDDLQPWHSFGADSVPANTE
						NEVE\PVDARPAADRGLTTRPG
		1			<u> </u>	NE VENT V DARFAADROL I TRPO

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	į.	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19080	49448	A	19189	1	2355	MKSIFTSEISSVLIFVNPEPGRVV
						SFLGSGRREHAEERARGPRETA
						AVAARAEQGRGGSHSHSSALG
						APRRVAMLPGLALLLAAWTV
						WAL\EVPTDGNAGLLAEPQIAM
1						FCGRLNMHMNVQNVKWDSDP
			İ			S\GTKTCIDTKEGILQYCQEVYP
						ELQITNVVEANQPVTIQNWCKR
						GRK\QCKTHPHFVIPYRCLVGEF
						VSDALL\VPDKCKFLHQERMGC
					1	FAETHLHWHT\AAKETLQ*RRS
						TNLHDYGMLLPCGIDKLRGVEF
						VCCPLAEESDNVDSADAEEDDS
						DVWWGGADTDYADGSEDKVV
						EVAEEEEVAEVEEEADDDED
						DEDGDEVEEEAKEPYEEATERT
						TSIATTTTTTTESVEEVVRVPTT
						AASTPDAVDKYLETPGDENEH
						AHFQKAKERLEAKHRERMSQV
}						MREWEEAERQAKN\LPKADKK
						AVF\QHFP\EKVESLEQEAANER
						QQLVETHMARVEAMLNDPRRL
1						ALE\NYI\TALQAV\PPRPRHVF\
						NMLKKYVRARTEGLVSTSL*HF
ļ						EHVR\MVDPKKAAQIRSQVM\T
						HLRVIYERMNQSLS\LLYNVPA
F						VAEEIQDEVDELLQKDANYS**
						RLWPTMISEPRI\SYG\NDAL\MP
			·			SLTETKTTRGSSVPVNEE\FSLD\
						DLEPVHFFWWLTSVPANTENE
						VE\PVDARPAADRGLTTR\PGSG
						LTNIKTEEISE\VKMDAEFRHDS
19081	49449	Α	19190	103	463	PLLYAMSTGKRWLSFGSMIPPV
						AICASNKRIPWPFSMSSLFMAW
						NKEMMFPVSGSLYPVTARSAD
						RSFSLNSMPTRPLL*ASSTGNLK
		1				LLTGSTIKPVAAWASSNSRAVP
						GRTKPIFMAS
19082	49450	A	19191	1	182	
19083	49451	Α	19192	3	437	MVHAEISDEGFFLSAPIALTQA
						WLNIPKT\RRTFCK\KCGKHQPH
						K\VTQYKKGKDS\LYAQGKRRY
						DRKQSGYGGQN*KPIFRKKAKT
						TKKIVLKALSALSPTCRSKSKC
						WLFKR\CKHF*TGEEIRKRKGQ
						VDPVLKCHLLFMEGH
19084	49452	С	19193	341	421	
17004	17772	<u> </u>	1,7,7,7	- ' '	1	<u> </u>

NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19085	49453	Α	19194	15		VLACTVEASVIGGQKREMNMG SIVREGFMEEAGPRIGFNLSFFI QSTVTGSSSVGSTQLQPLSLPPA TAGQEPRGTGPLGSHNSCPAPE PVSYGRLNDPPATTHPGSFTRF HSKPLTSSCWDSEMEGCVSNL MVCNLASFLNVSTNPCAFFPW YFYYLLLL*AGWSPLHIAASAG RDEIVKALLGKGGVLNVYTNT DRFLHCFLLSSPSFQIAVMLLEG GANPDAKDHYEATAMHRAAA KGNLLARNPNHYIELVVINWSL VVPDFRHLACDEERVEEAKLL VSQGASIYIENKEEKTPLQV\AK GGLGF*YSREWVGRFKQLGFIL YFVLLCLLSPVSYKLMYLCTRH
19086	49454	Α	19195	1597	1923	
19087	49455	A	19196	551	747	
19088	49456	Α	19197	1059	1296	
19089	49457	В	19198	1	1281	
19090	49458	A	19199	3	565	LPGRLLLLNFRQRRQIDRPSMN DTRNLSRT\RKFHAPTRPTSEEN KWVIDVLSPAGRPTVP*DQKFR EK\LA\KMYQAPHPDV\ILVFGF\ RT\HFGGGKTTGFG\MIYDSPWI MQRKNEP\KHRLAR\HGLYEKK KDLKKATEKERQEPELKKVQG GLAKGPLLGAGQKEEMKCLAV ELEIGSQPKELKVLQ
19091	49459	Α	19200	2	476	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19092	49460	A	19201	<u> </u>	2448	MRGSILKECGSKERASQVDLEY
						SRGNTASFAMCFHDRARIMKG
						SQRSRKLTGGEGREEGERVENK
						DFLDLHGQIQVRLVVKTASLLS
						RVTYIAEVEGRETDAGWRNEL
						QVGRLDSGPDILVVSISTDHTLV
						DTHLEAVGGHSHEFPYFKNHL
						ACHVESELQRDRNGHRDAGWR
						NMRHGITDKEMQDSQGSSDNH
						VLGSLRSWSARAAPALPGLPEE
		İ		1		AFVEMSTEGGFGGTSRSDAQQS
				1		LKSFWLRPRFFITELTLRAWIHT
						EDNNCRTYVAMAITKFDQLDF
						LIDIVLRDELKPPKCQEEVLQSV
						TPAEPVQYYFTLAQQPTA\VQV
						QGQQQGQTTASSMTTMQPGQII
						IAQLQQGQTTPVTTQVGEGQQ
						VQIVQAQPQGQAQ*AQSGTGW
						TVQGQIQTLATSAQPITQTEVQ
Ì		ļ				QRQQ*FSQFTDGQQLYQIQQVS
						IPAGQ\DLAQ\PMFIQSANQPSD
,						GQAPRSHEVIAAAPAARRERER
						ELAGRGPFCPPPAGNVRALKSR
ļ						RCLPGERRERARKKKPEGGAA
						ASERGETDSHPGASGQGPRDSG
						RVPLGMWSRRGLGVSRAPLHL
		İ				LLGVSTSFPLPLTPLIPTTAVVN
						PLKPLKNRIKAPHRRMPLVPGG
					}	SQNLGTVRARISTRQSHAEPST
						DNERVLSMTQDIHSIFLNLKNIT
						WICTLQSCGFLLSNLCKDIVHK
						AGSGEARTRWSPRAQAGTHED
19093	49461	C	19202	124	369	
19094	49462	В	19203	739	870	
19095	49463	С	19204	787	969	
19096	49464	A	19205	101	331	
19097	49465	A	19206	137	415	
19098	49466	A	19207	86	221	
19099	49467	A	19208	1	1347	FGARGNLKKAYSTRSKMAELN
19100	49468	A	19209	120	899	THVNVKEKIYAVRSVVPNKSN
•		1				NEIVLVLQQFDFNVDKAVQAF
		1				VDGSAIQVLKEWNMTGKKKN
		1				NKRKRSKSKQHQGNK\DAKDK
						VERPD\ACPLQPQPPQIQNGPM
			l			NGCEKDSSSTDSANEKPALIPRE
						KKISILEEPSKALRGVTGPNIEK
					[SAKDLQRCTVSLTRYRVMIKEE
						VDSSVKKITAAFAELHNCIIDKE
						ASLMAEMDKGIEEAMEILTAR
L	L		<u> </u>	l		Q\RKAEALKRLTDL\AS\QMAE

SEQ ID NO:	SEQ ID NO: of peptide sequence	I .	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19101	49469	A	19210	211	2150	ADTKPERGVSSAVFASGSEGRR LGCVLLSSSETRLLSGTLLWIPR AYSTRSKMAELNTHVNVKEKI YAARSVVPNKSNNEIVLVLQQF DFNVDKAVQAFVDGSAIQVLK EWNMTGNKKNYKRIRSKSKQH QGNKDAKDKVERPEAGPLQPQ PPQIQNGPMNGCEKDSSSTDSA N\EKPALIPREKKISILEEPSKAL RG\VTEGNRLLQQKLSLGG\NP KPIHGTTERSDGLQWSAEQPCN PSKPKAKTSPVKSNTPAAHLEI KPDEL\AKKRGPNIEKSS/VKDL QRL/CTVFF*LEYRVMN*RERK WDSSGERRSKLPFAE\LHNCIIE QKEV\SLMGRKWD*SLKEGRP WEILDWLVRKKSKKKLKRLTD LGQSRMAEMQLGPKLR\AEIKH FVSEAVNYDE\ELGKSCPGFSC DIEQ\LK\AQIIALPGKITHPK\NN YFLKELPCSSLL\PLLNAARSKP LGKTEVTFSRKSSTHNKPSEGK AANPKMVSSLPSTADPSHQT\M PANKQNGSSNQRRRFNPQ\YHN \NRLN\GPAKSQGSGNEAEPLGK G\NSRHGTQEGQPH/NRFGFRPK NKGRCQKIQEASLGIEGPPRAP AHFLKRPRAKGSHAADTFGRA RAFSGVS/GSVRGSQCNLFPT\RI EVSTDAAVLSVPGCDRWVA
19102	49470	Α	19211	93	180	
19103	49471	Α	19212	3	39	
19104	49472	В	19213	1	888	
19105	49473	A	19214	863	1509	GCAGPCLVNQMFTSSILCKSHC HSLVSINQGHNAPWKAAG\PLP LKAGYC\QSFSPCDSLKYG\SWD EKDLTVPQPDTRKASVLRWISQ RGKPLAVDMEEG\HCL\CLPPG N*NVLGCKTPIVHLFNSELGRK AALWREARHVG\SNAALLFFTP LRCLGGEKHKSGLHAHPGIVP\S LELNYDIDSFAHVFVFVVAVEL PLIITLLPYCIPLCCNNEK
19106	49474	С	19215	289	510	
19107	49475	В	19216	534	1686	
19108	49476	В	19217	l	1062	
19109	49477	В	19218	1	1878	
19110	49478	В	19219	1	747	
19111	49479	В	19220	230	786	
19112	49480	В	19221	67	6011	
19113	49481	В	19222	1	3455	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19114	49482	В	19223	1	4763	
19115	49483	В	19224	65	4534	
19116	49484	В	19225	512	4782	
19117	49485	В	19226	345	2194	
19118	49486	В	19227	267	7391	
19119	49487	Α	19228	578	739	
19120	49488	A	19229	419	1410	KQRQARERHLPRIPQLSAVVTIE TL*RK*RNFAELKIS/RLRKKFA HKMLQKARRKLIYEKVKHCHK EYRQMYRTEIQMATMARKAG NFFVHPKPKLAFVIRISAISGVSP KWKNKQLCPLNHVFGLGMGD LRAFGLADGTGPSGGEITGMRD LRGPLSVRAPSAGVAEIPRHLIV FEDVVNGKFLQTSRCEISGSKN KVWQTLLKNKRERQRLPTGFG RLADQRAYGEAKNNCGQVGL VETLWERHLLRIGVYIATSSLA VPTLSSNYNGLAHDQSCSDTVQ RSTGAPEAWDRGYCVGRGAQG IPIFFLDRPLTSLFDEARGSSSAS
19121	49489	A	19230	3	307	AGTMEGVEEKKKEVPAVPETL KKKRRNFAELKIKRLRKKFAQ KMLRKARRKLIYEKAKHYHKE YRQMYRTEIRMARMARKAGN FYVPAEPK\LAFVIRIRGI
19122	49490	A	19231	3	160	GRVLL*DKTSRGTIRQQQLLFT NICCSAASA/G*YPGKQGLEWT SSRLQQTCS
19123	49491	A	19232	303	570	GLCIRHVLVPSAAGLLEFAGGP LQTLFAWVSAAEAAEQQIWVN SKCC/CPDRSSGSFVSEEYLAV* GVGLTLSSASLMLGAVDWSCS HSA
19124	49492	A	19233	1	176	MQYVQQKPKLKFQVHICAYRK TVLQPPLLIPRQT/WVWSGPPA NSNRPAAGGADC*KEN

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
1				Sequence		
19125	49493	Α	19234	1	1977	MFAVLQAPLVIPRQT/WVWSGP
ľ						PANSNRPAAEGPDC*KEN*RQT
		l				GSGVDLQQTPTDLQLRVLIVRR
						KTNEQKGHPHQNPICTSPSSKT
						KVDTTTNMGEKQSRKTENSKN
		l				QSASPPPNLKKKREKNQIDAIK
						NDKGDITTDPTEIQTTIREYYKH
		İ				LYSNKLENLEEMGKFLDTYNLP
						GLNQEEVESLNRPITGSEIEVIIN
		ŀ				SLPTKKSPGPDGFTAEFYQRTK
						DKNHMIISIDEEKAFDKIQQPFM
						LKTLNKLGIDGTYLKIIRAIYDK
						PTANIILNGQKLEAFPLKAGTR
						QGCPLSPLLFNIVLEVLAREIRQ
						EKEIKGIQVGKEEVKLYLFADD
						MIVYLENPIISAQNLLKLISNFSK
			•			VSGYKINVQKSQAFLYTSNRQT
						ESQIMSELPVTIASKRIKYLGIQL
						TRGVKNLFKENYKPLLNKIKED
						TNKWKNIPCSWIGRINIMKMAI
						VPKVIYRFNAIPIKLPMTFFTEL
						EKTTLKCIWNQKRAHIVKSILS
						QKNKAGGIMLPDFKLYYKATV
						TKTAWYWYQNRHIDQWNRTE
						PSEIMPHIYSYLIFDKPDKTRNG
						ERIPYLIRWCWCWQNWLAICR
						KLKLDPFLTSYTKINSRWIKDL
į						NVRPKTIKTLEENLGITIQDIGM
						GKDFMSKTPKAMATKAKIDK
						WDLIKLKSFCTAKETTIRVNRQ
		İ				PTEWENISAIYSSDKGLISRIY
19126	49494	A	19235	206	456	
19127	49495	Α	19236	2	385	NLCCSAASAGDTQANRVWSGT
			1			PTNSSRPEVSSGAMDGEGWGL
						PVDPLTPGHQDALPWQRCYHP
						CSSSSVPPRQACASPASCSSSAA
						W/TSASTGPWHSGCGSSCGSCC
						CWGSPSASVGVGAGAIRSRTV
19128	49496	В	19237	1	642	
19129	49497	Ā	19238	128	584	
19130	49498	A	19239	4054	4578	QSGPSAAGLLEFAGGPLQTLFA
						WVSAAEAAEQRILVNSRCCCLI
			}			VP\RKFCLRGVASHVRCQSAPY
1						DVFCHGITLAKCWCSALGLPSL
						QSCEPNALFFINYPVCGI/TAPCC
1						KVEGTMG\VGPKSPQL\SYNVL
1						GSDNNGVLLGSMAIFTILILLIH
						EHGMFFHLFVSSFISLSSGL
	<u> </u>	L	l	<u> </u>	L	PITOINI LIIDI ADOLIDEGOOD

SEQ ID NO:	SEQ ID NO of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion
19131	49499	A	19240	11	948	MVLEKAREEVDPPSMLIPPVT
.,		`				CGQLPRSQCPQPTLSRMLTLE
						GISLGIIVLMISLWPDTVKEYO
						REALSFKKYNKNPGLLKFAG
						LQTLFAWVSPAEAAEQQISAI
						FLWKLRPRGAPTRCQPLLSYT
						CLSAPAGSFVYTMRGKPPIQA
						VMADASPPTKIEHPRTTSDCC
						ASKNFKPVDLSLLCSMGVGS.
						LDHWATWLQSPFQGSERFCL
						GVPGTTGV*KETPAASSVSAC
						AAQFCA*N/HRTLVV*APKGI
	1	1				WVCGLRRPWEKRSICAGVH
						SWHSPSWLPLARGGSSSTPCT
						WVRRCPTLHL
19132	49500	A	19241	171	1345	YCPKEGTLDFNFAFNMTAVN
19152	149300		17271		1343	ALIRDTKWLTLEVCREFQRG
						SRADADCKFAHPPRVCHVEN
		ŀ				RVVACFDSLKSVTAVPFLFM
						GLSREPETVPASMVQTFSLGV
					:	GGAGSV*PDENCKYLHPPPH
				-		TQLEINGRNNLIQQKTAAAM
						QQMQLMLQNAQMSSLGSFPI
						PSIPANPPM/SFESLRYHILGM
						LVPAELVPNTPVLIPGKPTSLO
						PGSCLGPKLMAFQINWEVLP
						FQRG\NC\SRGEND\CRY\AHP
						ASMIE\ASDNTVTICMGYIKG
						TVGVWGIKAKSSKQRILDTK
						HQGNSFVRGTDYLIPICTNMS
						AKGTAFLKWMQKLVYNGSS
						OCSPAFFCCAWKEHSEKAGR
						SGAKCNRHRHTNPVFLMREC
						IS
19133	49501	A	19242	61	544	KFNP\FVTSRPRARIRKRAF\N
17133	17501	1	17242			\SHIPKGRLLSFPPLSPKSLRPE
						TNRADSHAPIPKRMIESSRLY
						GHL*GSAKLAKVVPRFTRKK
						VIYIER\VQREKA\NGTTVH\V
		}				HPSKQVVITKAKTGTKTRKK
	}		2			PNRKAQSRQVGKEKG\KYKE
						TIEKMQE
19134	49502	A	19243	652	1042	TPLSSCFLSREKATQLLFPAV
19134	77302		1,7243		1072	VPMVLSAMGFTAAGIASSSIA
						KMMSAAAIANGGGVASGSL
						TLQS/LG*VSWRGLPTVLPTE
						RQACPTLSTLCFFPGATGLSG
						KFILGSIGSAIAAVIARFY
19135	49503	A	19244	48	397	KI ILUSIUSAIAA VIAINI I

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19136	49504	Α	19245	42	296	GCVGVRPSLHPATSTASGSASP
						TLARAMASVSELACIYSALILH
						DDEVTVTEDKINA/APAEEKKV
						EAKKEESEESDDDMGFGLFD
19137	49505	Α	19246	172	519	LGPSFLHDDEVTVTEDKIQMPLI
						KAAGVNVEPFWPG/LCFAKAL
				İ		GKRSTLGSLILAMLGGPVGPGS
		Į.				QPAGCLQPAG\GSCPPSTCCWF
	İ	Ì				QLEGEEKWEAKERKIPREF*LID
						MGFGLFD
19138	49506	Α	19247	3	175	
19139	49507	Α	19248	2	390	GWDWNCVWEPHHWLCQSL/N
						SVTQAGVQLCNLSSLQPLPLGF
						KQFSCLSLPSSWDYRNPSLKQQ
						LFSYAILGFALSEAMGLFCLMV
						AFLI\SLPCEGAVSTSHSSPASG
		l .				WPRVFLFLYLPRQPGERGWLR
19140	49508	В	19249	139	13320	
19141	49509	Α	19250	28	450	TNDFNLQNPGSCFPSACPAGVG
						SERLPVLGWFVGNRSFPPPPQST
						LSVLGKRLGRNGAIAAGVFGA
						LFIGYCIYFDRKRLK*PQLKDAE
1			ł			AVQKFFLEEI*L\GEEILAKGVD
]				İ		HLTNPSAFFG/QPNHFPLQMDSF
		<u> </u>				GPFLRFQA
19142	49510	Α	19251	164	420	OTHER WILLIAM OF THE WILL
19143	49511	Α	19252	1	329	STHTTY WEGCRTTSAGLLRKH
						EPGEEAF/FYHST\SKKPGIAGLI
						KVNVQFVRMMKRSIPWAELNP
	-	E				YHQAHKATGGPLKNMVLFTRQ
10144	40512	D	19253	39	368	RLSIQPLTQEEFDFVLSLEEKEP
19144 19145	49512 49513	B A	19254	1014	1290	
19146	49514	A	19255	122	680	LARITRIVRTKVPCSVTMSRPRK
19140	47714	Γ	19233	122	080	RLAGTSGSDKGLSGKRTKTENS
						GEALAKVEDSNPQKTSATKNC
						LKNLSSHWLMKSEPESRLEKG
						VDVKFSIEDLTAQAKQTTCWD
						GVRNYQAS/RNFLRAMKLGEE\
						AFFYP*QLAKKPGIAGLMKIVK
	[EAYPDHTQF*EK\NPHYDPSSKE
						DNPKWSMKSLILF
L	1	<u> </u>	<u> </u>	<u> </u>	<u> </u>	DIALKASIMIKSTIFL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hođ	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19147	49515	Α	19256	2	1005	RISTSTLCPKPKKLIAPPGRAGL
						ASMGLNEEQKEFQKVAFDFAA
						REMAPNMAEWDQKDRKADGK
						GGAVTQEVQAQLLVKHGQKL
						VSRFQLSSFSYVDSVRTTQSKQ
						EHLQTDKGLSGKRTKTENSGE
						ALAKVEDSNPQKTSATKNCLK
						NLSSHWLMKSEPESRLEKGVD
						VKFSIEDLK\AQPQ\QTTCW\EG
				:		VRNY\QARNFLRAMKLGEE\AF
						FYH\SNCK\EPGIAGLMKIVK\EA
						\YPDHTQFEKKQPPPL*PPS*QR
						GTTPKWPHGWMV/QVWFGMM
						KRFHSPLAE\LKS\HHQAHKA\S
						GGPL\KNMVLFTR\QRLSI\QPLT
				· . <u>.</u>		\QEEFDFVLS\LEEKEPS
19148	49516	Α	19257	3	727	VRLQFLTPTLRAARTMAAPPQL
		1				RALLVVVNALLRKRRYHAALA
		1				VLKGFRNGAVYGAKIRGPSRR
		į				WVMTFLFREWAASREKLWAIL
						Q\ATYIHSW\NL\ARFVFHLTRV
		l				FRCPAVPYITRARTYP\AHAFPG
						AGLSFGGIL\VFGRKQ*TFNSQI\
						NMYLLSTRPCLP*SR\LAVRRK
						GLHP*NPRWDPFP/VCLTAVVW
		1				GLVLWLFEYHRSTLQPSLQSSM
						TYLYEDSNVWHDISDFLVYNK
	10515	ļ. —	100.50		0.40	SRPSN
19149	49517	Α	19258	1	849	MSGALDVLQMKEEDVLKFHA
		1				AGTHLGGTNLDFQMEQYIYKR
		1				KSDGIYIINLKRTWEKFLLAAR
		1			1	AVVAIENPADVSVISSGNTGQR
		1				AVLKFAAATGATPIAGHFTPGT
		1				FTNQIQAAFREPRLLVVTDPRA DHQPLTESSYVNLPTIALCNTDS
		1				PLRYVD/ICNNKGAHS\DPEEIE
		1				KEEQAAAEKAVTKEEFQGEWT
						APAPEFTATQPEVADLSEGVQV
						PSVPIQQFPTDDWSTQPATENW
						SAAPTAQATEWKMQQQSTILE AESSPHQTPNLPAPWSRTSQPPE
		<u>L_</u>		<u> </u>		MESSPHQ I FINLPAP WSK I SQPPE

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19150	49518	A	19259		1309	GNLAAGKTIQAQDRDAVGILSS RTGESMENLQKNLLPKQRRRT RETFTMSGALDVLQMKEEDVL KFLAA\GTHLGGTQFLLSRGHG YFYKGKS/DGLSFIINPKRAREE\ LLLPARQIVPHKTPS\DVSVISSG NTGQVCGTVRA/VCLKFAAAT G\ATP\IAGR\FTPGTFH*PRSKA AF\REPR\LLVVT*PPGADPPALS RSASYVNLPYHLRCGNT\DS\QL RYVDIAIPCNNKG\AHSVGLMW WMLARGSFCAMRGTI\SREHPW \EVMPDLYFYRDPEEIEKEEQA AA*E/RQVTKEEFQGEWTAPAP EF\TATQPEVADWSEGVQGAPL VPYSAIPPTEDWSSQPAMEDWS AAPTAQATEWCSRTSLRQNPSD TELKKEGLYSARSISKTHVSKT ELPNQGTKKCSRTSHRQNPSVT
19151	49519	A	19260	965	2495	ELKKEGLYLARSIGKTHISKTEL AFTTRSTFSTNY\RSLGSVQAPS YGARPRQAARPASYAGAGGSG SRISVSRSTSFRGG\MGSGGLAT GTSPGVMAGMG\GIQNEKETM QS\LNDRL\ASYLGQE*RSLEDR RTRKLGRAKFREHFGRKKGPQ V\RDW\SH\YFKII\EDLEGFRFFR KILVGQCPASVSARFDNAPSLL LD*F*EFKY*GQSWPMCPVLLE NDIPLGSAKVI\DDTNYHTDLQL \ETEIRGSSRRELLFQ*RRNHEE\ EVKGLQAQIASSG\LTVE\VD\AP KSQDLAK\IMADIR\AQYD\ELA RKEPEKEA*TKYWSSADLREST TVGHPHKSA*GWKLAEDDASQ KLRRT\VQSLEDRPWTSMRKSE RPALENSP*GEVEAPLRPLQDW SQLQTGIPACHLWSQELGTRHR AQRGQRPGPQGV*GPLLNHQG SSLEAEICHLTRPPSWKIGEDFN L\GDS\LDERNSMQTIQKTTTPP G*VGLGKVVS*DQLTPKVLKAL SQQEAVVPFWGSRRPIKSFRVQ KKKKEKESF

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19152	49520	A	19261	387	1125	CCRPCATPYALRSPLR*PACPPP
						RWRPSSMPTSRPPVSPAASPTC
						GTCMLSP\CPPAEHLLHLPHAQP
						QDPALLQPCHPAPPLRPRALHA
						AHHVGWPAPHQPLLPPPVLCPC
						GGAGRGGG*RCPWRGPSSSSPG
		1				PTLIAGQDPGAAQP*TWPQ/PTL
İ			:			TSVSATASPKAP\LPLAAAVPGA
		1			:	PAEGCCPLLRQALPAGQLPPGG
						RRATSAVPPAVPGD/PPAQPTM
		1				PGSMSCWALAPARPCPPGRCW
]				AWRS
19153	49521	Α	19262	3	1097	RGNSRLRYSHEDELQLPRLPEL
		1				FGNCRQLLEEVEVAT\DPAGFR\
	8	1				IVQEK\VFKG\LDLLE\RAAEML\
						SQSTLFGRECRFWKDLSPTHLIT
		ŀ				FWGP\AFQGALTM\KQVNPSKR
						LRSFCSGAREHFI\NYLT\QCH\C
						YHVGQSFEL\PKT\MNNSAENH
						TANSSMAYPSLVAM\ASQRQA
						KIQRYKQKKELEHRLSAMKSA
						VESGQADDERVRGYYLLHLQR
	1					W\VDISLEEIESIDQEIKILRERDS
8						SREASTSN\SSRQERPPVKPFHS
						HFGTWLHSQSTFGAGYSKGWP
						TYGR*VSWYE\QHRKLWSTYP
				1		DQGNMPRAAPEGISEKAAQQQ
						EDQEEKEEEDDEPNNFHRARE
]		W\DDWK\DTHP\RG\YGNRQNM
19154	49522	Α	19263	2	262	
19155	49523	Α	19264	3	1062	GRTAENPARAVSSPNFYAHRKT
						EVLFPCLTQPLARGPKKHLK\R
						VAAPKH\WMLDKLTRVFAPR\P
		ł				STG\PHKLR\ECLSRIIF\LRNRLK
		l				\YAL\TGDEVKKI\CMQR\FIK\ID
		ŀ				GQGSELDINLPLLGFMGCPSAF
:	ŀ					DKTGEENF\RLDLIDTQGVRFC
		1				WYIRI*PPWRKAKLQVWCQK*
		1				GKILCGPTKRESPSSGLTH*CPA
		l				PHPAYPPIPPHPR*NGYPIPD*I*E
						DLAKINLISIQSSEHW*TLCMGD
		1				LEGA*P*GRNWVLITPTERGHP
						GIFLTVGSR*KDANGQQAFGHF
						GLFQHFLVIGKGNKTHGISL\PR
						GKGIRLHHLLEERDKRLA\AKQ
						SQWGEMGSLGVDIVQIFWYRN
19156	49524	Α	19265	62	298	

		SEQ ID NO:	I .	4	Amino acid sequence (X=Unknown,
NO: of peptide sequence	hod	in USSN 09/540,217	location of first	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
Sequence		0772710,217	sequence	or pepilae sequence	derendi, v possiole nacicolide insertion)
			<u></u>		
19157 49525	Α	19266	1	555	MAWQMMQLLLLALVTAAGSA
					QPRSARARTDLLNVCMNAKHH
					KTQPSPEDELYGQ/SP/WMCKG
					SCRKTKSWNI/HRKSKCEVGLA/
					WEACSVSAGTGRGPGCGRWV
					GAPQGP/CPRKCSSG*PTW/VQR
1					SQNMEEMAVVNQSWRKERILN
					VPLCKEDCERWWEDCRTSYTC
					KSNWHKGWNWTSAPSAVCDP
					LL
19158 49526	Α	19267	216	974	MDMAWQMMQLLLLALVTAA
					GSAQPR\SARA\RTDLLNVCMN
			:		AKHHKTQPSPEDE\LYGQCSP\
	ľ				WKK\NACCTAS\TSQELHKDTS
					RLYNFNW\DHCG*KWNPTCK\R
					PLYPQDSCL\YECLTPTLGPW\IR
					QFNQSWRKERFLNV\PLCKEDC
					ERWWEDCRTSYTCKSNWHKG
			ļ		WNWTS\GINKGRPGAF*STFES
					YFPTPAALCEGLWSHSFKVSNY
					SRGSGRCIQMWFDSAQGNPNE
					EVAKFYAAAMNAGAPSRGIIDS
19159 49527	Α	19268	345	2138	QRLTATSSWTTMAKNRRDRNS
	ŀ				WGGFSEKTYEWSSEEEEPVKK
					AGPVQVLIVKDDHSFELDETAL
					NRILLSEAVRDKEVVAVSVAG
					AFRKGKSFLMDFMLRYMYNQE
					SVDWVGDYNEPLTGFSWRGGS
					ERETTGIQIWSEIFLINKPDGKK
					VAVLLMDTQGTFDSQSTLRDS
					ATVFALRHNDPASIQVYNLSQN
					VQ\EDDLQ\HLQLFT\EYGQDW
1					AMGGKHFLEAIFRSLDYFFVRD
					W\SFPYEFSYGADGGAKFLEKR
					LKVSGNQHEELQNVRKHIHSCF
					TNISCFLLPHPGLK\VATNPNFD
					GK\LKEIDDEFHQKL*KYLIPWA
					YFRSRGA*DI*RRSNGGLKI\T\C
					RGLVEYFKAYIKIYQGEELPHP
					KSMLQATAEANNLAAVATAK
		1			DTYNKKMEEICGGDKPFLAPN
					DLQTKHLAT*GRNLWKLFRGV
					KKMGGEEFSRRYLQQLESEIDE
					LY\TNISSHNDSK\NTFPCSSYPQ
	1				PTLFCSHPLITYVIAGVTGFIGL
		1			DIIASLCNMIMGLTLITLCTWAY
					IRYSGEYRELGAVIGPRWLQLC
					GTREVQMRPWNKLYQCQQAT
					PTDILYPSKPFPLHPKVGNLTEQ
					IQKRKKMVMPKF

SEQ ID NO:	SEQ ID NO: of peptide	Met	SEQ ID NO:	Nucleotide location of first	ł .	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19160	49528	A	19269	351	509	VVPLPLPPLDSWTCESWLRSHI LTGESENPDHKRI*PYTSDLPEG
10161	10.550	↓	10070	701	4224	PGPWATD
19161	49529	Α	19270	781	4324	ESVREKVLCGLIQQTVTSVC/W
						SGTYIFAVLLVCVVFHSGAQEK NYTIREEIPENVLIGNLLKDLNL
					:	SLIPNKSLTTTMQFKLVYKTGD
						VPLIRIEEDTGEIFTTGARIDREK
]		LCAGIPRDEHCFYEVEVAILPDE
						IFRLVKIRFLIEDINDNAPLFPAT
	:			1		VINISIPENSAINSKYTLPAAVDP
	ļ					DVGINGVQNYELIKSQNIFGLD
						VIETPGGDKMPQLIVQKELDRE
						EKDTYVMKVKVEDGGFPQRSS
19162	49530	A	19271	3	644	FRSLPRRCLPRWQPSSFLGHHG
1,710-	.,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,					PPFRPPCEAPKIVKKENPRKFIR
						APSQTGY\AKLSVNW\RKTPEAI
						DN\RVR\RRF\KGQILMP\NIGYG
						SNKK\TKHM\LPSGFRKFLV\HN
						VKEPEIAADVATNLTVPRSLTM
						FPPRNRKAIRGKGACPIGPFRSS
						PNPIARVAQLKEN*VRQLMCN
						KSYCAEIAHNVSSKNRKAIVER
						AAQLAIRVTNPNARAQ
19163	49531	В	19272	27	161	
19164	49532	Α	19273	80	430	RWLATHYLTVKLSLLPNLAVT
						CQPKREQNITSNESTDC*VIVTS TKSDSLYTVGMLALSVRAIRCP
						LYLLTGLIFVSKNDLWYCELQS
						HYTVVLLLFMRKRSRFQSNVQ
	1					RKLMTNF
19165	49533	A	19274	482	832	RWLATHYLTVKLSLLPNLAVT
15105	1,7333		''	1.02	032	CQPKREQNITSNESTDC*VIVTS
						TKSDSLYTVGMLALSVRAIRCP
				:		LYLLTGLIFVSKNDLWYCELQS
		1				HYTVVLLLFMRKRSRFQSNVQ
						RKLMTNF
19166	49534	A	19275	121	201	SMYMDEN*RLELITL*FSSMYM
		l				DHSLL
19167	49535	В	19276	1	1147	
19168	49536	Α	19277	305	442	AGLFKSRKVSSICIRFE*ILTSCL
						GVRISTGKLVARIHGSFADAE
19169	49537	Α	19278	133	527	GFLERPRILIKVTLSIKQKSTSIF
						VGGQFSKFRFLTVLAWKCLSQP
						CF*A*TPASQHVS*SIRAISSMIS
						KRSELLVRLWRRPSP*AGLFKS
						RKVSSICIRFE*ILTSCLGVRIST
10150	10520	 	10020	521	660	GKLVARTHGSFADAE
19170	49538	A	19279	531	669	AGLFKSRKVSSICIRFE*ILTSCL
		ل	L	<u> </u>	L	GVRISTGKLVARTHGSFADAE

SEQ ID NO:	SEQ ID NO: of peptide	Met	SEQ ID NO:	Nucleotide location of first		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO:	sequence	nou	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19171	49539	Α	19280	80	428	RWLATPYLTVKLSLLPNLAVTC
				•		QPKREQNITSNESTDC*VIVTST
						KSDSLYTVGMLALSVRAIRCPL
						YLLTGLISVSKNGLWYCELQSH
1					1	YTVVLLLFMRKRSRFQSNVQR
		<u> </u>				KLMTNF
19172	49540	Α	19281	10	426	
19173	49541	Α	19282	1	900	
19174	49542	A	19283	314	451	AGLFKSRKVSSICIRFE*ILTSCL GVRISTGKLVARTHGSFADAE
19175	49543	Α.	19284	1	1139	MWGIGSRTPIYNKILVYSSPEVS
					}	CEKPKYVKSQTSTHTGFESHKR
					,	YIFYLRLVGKNLHVSAMGLGK
						SQERRKKALKSQPQQPADSFPS
		:				LPTVEGLTTSGEVAAGFWLSGE
						ALSSC*LLSASVHLLLACLGMG
		<u> </u>				KCSCSGFWLYF
19176	49544	A	19285	1	864	
19177	49545	Α	19286	306	444	AGLFKSRKVSSICIRFE*ILTSCL
	<u> </u>	ļ				GVRISTGKLVARTHGSFADAE
19178	49546	Α	19287	189	1089	HHTAHCQ*CWL*S/DHGINPLRS
						WVGTG*V/GVRGKVQYADLGA
1						ENWKPISNLHDMSSSHSKTLGY
						KRLTKSNPISCQILLYKSRSKGR
						KNQRSTRTHCHHPSPKIYSASA
		l				KEPWVLATNLPVEIRTPKQLVN IYSKRMQIEETFRDLKSPAYGL
		i				GLRHSRTSSSERFDIMLLIALML
						QLTCWLAGVHAQKQGWDKHF
						QANTVRNRNGSLLTNWPSVPP
						TIKEEENSEEELAAATTSKEQEP
						IGTOLDAVRTPEPLEEFPKREDQ
						EGSPPETSLPYKWVVEAANLLI
						PAVGSSLSEALDLIES
19179	49547	A	19288	1	549	
19180	49548	Α	19289	80	430	RWLATPYLTVKLSLLPNLAVTC
						QPKREQNITSNESTDC*VIVTST
						KSDSLYTVGMLALSVRAIRCPL
		1				YLLTGLISVSKNGLWYCELQSH
						YTVVLLLFMRKRSRFQSNVQR
						KLMTNF
19181	49549	Α	19290	80	430	RWLATHYLTVKLSLLPNLAVT
						CQPKREQNITSNESTDC*VIVTS
						TKSDSLYTVGMLALSVRAIRCP
						LYLLTGLIFVSKNDLWYCELQS
						HYTVVLLLFMRKRSRFQSNVQ
			L_			RKLMTNF
19182	49550	A	19291	773	1015	AGLFKSRKVSSICIRFE*ILTSCL
		1				GVRISTGKLVARTHGSFADASF
						STDLYHGTLKPASLTMSGVVLL
		1				GRMLARSARNWS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19183	49551	В	19292	1	1488	
19184	49552	A	19293	2	133	YAQTDDNRRNLILNGTPLRRVK VLVFDIEYP**KVYGYNSTLA
19185	49553	В	19294	1	879	
19186	49554	Α	19295	80	428	RWLATHYLTVKLSLLPNLAVT CQPKREQNITSNESTDC*VIVTS TKSDSLYTVGMLALSVRAIRCP LYLLTGLIFVSKNDLWYCELQS HYTVVLLLFMRKRSRFQSNVQ
			1000		1,700	RKLMTNF
19187	49555	В	19296	1	1575	
19188	49556	С	19297	1	2187	
19189	49557	A	19298	473	613	AGLFKSRKVSSICIRFE*ILTSCL GVRISTGKLVARIHGSFADADT
19190	49558	Α	19299	410	547	AGLFKSRKVSSICIRFE*ILTSCL GVRISTGKLVARIHGSFADAE
19191	49559	Α	19300	876	1005	
19192	49560	A	19301	167	263	KLIEAIWSS*FITGSSLRHNERLG ASTSSGEY
19193	49561	Α	19302	258	396	AGLFKSRKVSSICIRFE*ILTSCL GVRISTGKLVARTHGSFADAE
19194	49562	С	19303	1	1641	
19195	49563	Α	19304	1221	1327	ALHPRNRHLHKERLGCIPVAC* LYLFGQYDAHCTC
19196	49564	С	19305	1	1353	
19197	49565	В	19306	1	2047	
19198	49566	A	19307	80	429	RWLATHYLTVKLSLLPNLAVT CQPKREQNITSNESTDC*VIVTS TKSDSLYTVGMLALSVRAIRCP LYLLTGLIFVSKNDLWYCELQS HYTVVLLLFMRKRSRFQSNVQ RKLMTNF

1	1		l .		Amino acid sequence (X=Unknown,
of peptide	hod	in USSN	i .		*=Stop codon, /=possible nucleotide
sequence		09/540,217		or peptide sequence	deletion, \=possible nucleotide insertion)
49567	Α	19308	1	2597	MAMFMLLAAALLSTFLTNDVA
					LYTVVPLTITLKRLCEIPYNRLII
		i			FEGAGSQRWFATDANWQPAK
					YSYLGTFWSFVCRIYCPNGTAG
					WRNDADAPAPVLAMQYHTGV
		•			QTPEWKPRLVWSCLGLYIVFLT
					ALEFKQELWGLVIVAAGFALL
					ARRVVLSVDWTLLLVFMAMFI
				٠	DVHLLTQLPALQGVLGNVISNV
					PSTILLLNYVPPSLLLVWAVNV
					GGFGFLTCRHARIAAPADLKDG
					LAVHEKRETRSLTEKTGKMLLP
					KLGVGSRECGGNVKEVGIRAIE
					EFGPYKINGDKEIMRRMDDLLQ
					GFDAQHRMKLPAQPTSLATKS
					ARRVRYPLFCPSIFNGFFEEFIFH
					RFLAEQALEFFNLLHGGSKFRG
					RNNLFSGGDSSRKNQRSTRTHC
					HHPSPKIYSASAKEPWVLATNL
					PVEIRTPKQLVNIYSKRMQIEET
	ĺ				FRDLKSPAYGLGLRHSRTSSSE
					RFDIMLLIALMLQLTCWLAGV
					HAQKQGWDKHFQANTVRNRN
					CKMATLKEKLIAPVAEEEATVP
					NNKITVVGVGQVGMACAISILG
					KSRAMEHTTAGAPYTPRQQTC
					QCNMMSLADELALVDVLEDKL
					KGEMMDLQHGSLFLQTPKIVA
					DKDYSVTANSKIVVVTAGVRQ
					QEGESRLNLVQRNVNVFKFIIP
					QIVKYSPDCII\IVVSNPRDI\LTY
					GTWKLSGLPKHRVIGSGCNLDS
49568	A	19309	435	777	RWLATHYLTVKLSLLPNLAVT
	-				CQPKREQNITSNESTDC*VIVTS
					TKSDSLYTVGMLALSVRAIRCP
					CRLVDWADIPEQKQLMVLRAS
					VALHGRSVTLYEKAFPAFRAM
					FKRSS
	of peptide sequence	of peptide sequence hod sequence 49567 A	of peptide sequence hod in USSN 09/540,217	sequence 09/540,217 codon for peptide sequence 49567 A 19308 1	of peptide sequence hod grade se

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19201	49569	ĪA	19310	69	1270	LCLLRHDCYRGLQSLLLILCKM
		1				ATLKEKLIAPVAE\DEA\TVPNN
		1				KIT\VVGV\GQVGMA\CAISIPGK
						VSWPDEL\ALVGCFWEDKLKG
						GNDGSFQHGELIFFQTP*NCGQ
						DKGLFL*PANS*RFVSW*LAGS
						PVQPRRGRIRLQSWVPEKWLM
						VFQIPLFLKIVQVQSLICIIIGGFN
						PQLDILTYVT\WETKVGLPKHR
		ŀ				VIGSGCN\LDS\ARFRYLMAEK
						TWHFIPSSC\HGWNFGGNMGDS
						SV\AVWNGV\NVAGVSL\QEL\N
l						
l		ŀ				PENGNLTNDKLKIWKEGA*IW
l						WLKSA\YEVI\KLKGYPQLGAI
l						WI*VLAESYWNPMLKKSYPRI\
İ						HPGCPTNG*RQGMYGHWRME
		1				SPSLEPSLCIPQCPRGLNPAVIQP
						RSLKDDEVASASRKSARIPLWD
		1				NPEWTLKRPLT
19202	49570	Α	19311	52	406	HVEAAGSR*CAAPAGALLRFLG
						TPSFEDVSIPEKPKLRFIERAPLV
						PKVRREPKTLSDIRGPSTEATEF
į				İ		TEGNFAILALGGGYLHWGHFE
						MMRLTINRSM\DPKNMFAIWR
						VPAPFQAH
19203	49571	A	19312	1	783	
19204	49572	Α	19313	15	397	LLAEWTMGHTPVGGFRGAISL
				Ī		DAAP*EQGTMLVSWLCDLFLV
				İ		LRILVGKTVSSKNIFIERVNGDG
				ľ		KHENIPMSSPPRSWFPSEKRFTN
						RRKTSSHLQNLVLGAVSLRQH
						WRGGVPRNLTRAPAGAAH
19205	49573	В	19314	127	181	
19206	49574	Α	19315	242	1067	CAAPCGCPCQIPGALLPASAGV
ĺ						KTLLPVPSFEDVSIPEKPKA*DL
ĺ						LERAPTCAQK*RREP*KFKVTIR
ĺ						GP\SH*KPTEFYRKAIFAILGIGV
ĺ						VGYLHWG\HF*K*WRLTIQPAL
ĺ			1			WTPRTWFAIWR\VPAPFKPITRK
ĺ		1		İ		SVGASHGGEAKGAI*HYLTP\V
İ	}					KAGRLV\VEMGGRCEFARSAK
ĺ						VSLTQVALRSCPFASKRL*ACG
						TPREGCEQDQEERERQQPRTPW
ĺ						TI*AE*PTAQHGWAYGKVPEPH
	-					MTLTPQGEILGARFYMPKTCVV
i	I	ı	1	1	1	1
Į		-	į.	i		SVGDILYIGY

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	1	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19208	49576	A	19317	104	763	SAAADGPDRLRRRPPANLRARP
19200	7,570	^	17517	1104	1703	VLAPASATASARRACSPPHPQT
						MSESK\SCPEYASFFAVMGASA
						A\MVFSALGAAYGTAK\SGTGI
						AAMSVMRPEQIMKSIIPVVM\A
						GIIAMLRPGWVAVLIANSLNDD
						ISLYKS\FLQLGAGLSVGLSGLA
						QPAFA\IGFVG\DAGVR\GTAMH
						AADIFRGA*FLILIF\AEVLRL\Y
						GLIV\ALILFHKVDLFRAHQPTE
19209	49577	A	19321	36	164	GET V VIEWET THE V EEK TO THE TE
19210	49578	A	19322	190	743	NDRGKEGSLSTVPSIRNTDSGH
17210	17270	`	17322	150	, 13	LAPRRKPQLLKPSLNTSFPKAR
			ĺ			KIHKGEHEVASAGGNPPAPILP
İ						KVPP*FNIHRRGPPWPDHVRFG
				1		SLF\NTLFL\NW\CCLGVSLAFRL
						LP*KSRDTGKMGLGDVEPGAQ
						ALCLPPPKCLNI\WALILG\IL\MT
						IGFIL\LLVFGSVDSLTILWFQIIQ
						GKTGGY
19211	49579	A	19323	2	257	SK1001
19212	49580	A	19324	 	1227	MGRARKSGLGQRSRPTASRSEA
19212	17,500		17324]	1227	AVQPGVRKARGAGNWRVGLQ
				}		TGEAAPSPHRDLRDTPDPRPWL
						ARTHRMTTTLVSATIFDLSEVL
ľ						CKGNKMLNYSAPSAGGCLLDR
:						KAVGTPAGGGFPRRHSVTLPSS
			İ			KFHQNQLLSSLKGEPAPALSSR
						DSRFRDRSFSEGGERLLPTQKQ
						PGGGQVNSSRYKT\ELCRPFEE
		1				NGACKYGDKCQFAHGIHEL\RR
1			1			LTLHTKD*TELCRTFHTIGFCPY
						GPRCHFIHNAEERRALAGARDL
						SADRPRLQHSFSFAGFPSAAAT
						AAATGLLDSPTSITPPP\ILSADD
		ł				LLGSPTLPDGTNNPFAFSSQELA
		1				SLFAPSMGLPGGGSPTTFLFRP
		1				MSESPHMFDSPPSPQDSLSD\QE
		1				GYLSSSSSSHSGSDSPTLDNSRR
						LPIFSRLSISDD
19213	49581	A	19325	204	456	
19214	49582	A	19326	1	323	IVEYHGPGVDSISCTGMATICN
17214	17.502	` `	17520	ľ		MGAEIGATTSVFPYNHRMKKY
		1				LSKTGREDIANLADEFK\DHLVP
						DPGCHYDQLIEINLSEMV*PDM
						LPAPRLAHGVTVVVVGGFLK
19215	49583	Α	19327	2	131	E. H. C. L. L. L. L. L. L. L. L. L. L. L. L. L.
19216	49584	C	19328	2	283	
19217	49585	A	19329	1	206	
19217	49586	В	19330	217	435	
17210	77300	12	17550	1217	1.55	L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19219	49587	Α	19331	1	565	PWE*SPKGIGVSCGPLSGWS/SP
						KD/VILKVQ/GILRFKGGTVQIV
						ESTGLG*TPFSCTGMRQFCNMG
						AEIGATTSV\FPYNHRMKKYLS
		ŀ				KTGREDIANLADEFKDHLVPDP
			:			GCHYDQLIEINLSELKPHINGPF
			:			TPDLAHPVAEVGKVAEKEGWP
						LDIRVDALACQSLAFGAAWVQ
	l					CGCEEHGAWSQMAWA
19220	49588	Α	19332	117	2603	GQHCLLPWGVFPFLKKALGV\R
		i				Q*\HVASVLCQRAR\VAMSHFE
						PNEYIHYDLLEKNINIVRKRLNR
		·				PL\TLSE\KIVYGHLDDPASQEIE
						RGKSYLRLRP\DRVAMQDATA
						Q\MAMLQFISSGLSKVAVP\STI
					}	HCDHLK\EPQVGGEKDL\RRAK
		l				DINQEVYNFLATAGAKYGVGF
						WKPGSGIIHQIILENYCVPWLSF
			E			*LGPDFHTPNGGG\LGGI\CIGSC
						GWPNAWEWNGLGIPWE\LKCP\
						QVIGVKLTGSLSGWSSPKDVIL
						KVAGILTVKGGTGAIVEYHGPG
						VDSISCTGMATICNMGAEIGAT
						TSVFPYNHRMKKYLSKTGRE\D
						IANLA\DEFKVHLVP\DPG\CHY
						EPTELKINLSEL\KPHIQWGPFTP
						DRGSPCWQEVGKVGREGKGW
						PL\DIRGGS*LVACTNFKLWKIW
						GRSAA\VAKQALAHG\LKCKSQ
						F\TITPGFPRQI\RATFE\RDGYAQ
						ILRDLGG\IVL\ANACGPCIGQW
						D\RKDI\KMGEKNTI\VTSYNRN
						FTGRN\DPNPETHAFVTSPEIVT
		1				ALAIAGTLKF\NPETD*PDGAR
						MGKKFQAWRFPD\AD*VFPKG
						EF*PRGRDTYQHPPK\DSSG\QH
						VGT*APRSQRLQLLDPFDKWD
						GKDLEDLQILIKVKGKCTTDHIS
						AAGPWLK\FRGHLDNIPN/NHLL
						IGAINIENGKANSVG\NAVTQEF
						GPAP*QLAR\YYQETLASRLGW
19221	49589	В	19333	1	1311	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	1	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19222	49590	Α	19334	2	1390	PRVRPRVRARPTKMSALGSLSL
	ŀ	1				PSVLWGQLF\AVDSGNDVTDIA
ŀ						DD\GCPKPPEIAHGYV\EHSVRY
						QCKNYYKLRTEGDGVYTL/NTI
						EKQWIKLRLFGD\KLP*IVEADD
:						GCPKPPEIAHGYVEHSVRYQCK
						NYYKLRTEGDGVYTLNNEKQ
						WINKAVGDKLPECEAVCG\KPK
						NP\ANPV\QRDPGVGHL\DCQRS
						FFPWARLKDGFPHH*SSTHQGP
		l				RWIQLNQWLA*PRLKNLLSLN
						H\SENCQQRKDICPYF*HFMWG
						KKQLVGD*GRLFLQP*LTPRVD
						IGLI\KLQTEGCSC*LRRVMPICL
						PSKGLWQK*GRVGLCFLAGGR
						NAQF*NLLTHLKVCPCCPVA*P
						KTQCHKAIYERPSNKSPRKRKN
						TRKKPL*GVQAPILK*NTPFLCL
						AMS*RYPKERHPAYG/QNAGAS
						ALCPFHRPLEEDTPGMPTGIL\S
		ŀ				FDKS\CAVA\EYGVYV\KVTSIQ\
						DWVQKTIA
19223	49591	Α	19335	65	665	GVRGFWAGTMASRAGPRAAG
						TDGSDFQHRERVAMHYQMSVT
						LKYEIKKLIYVHLVIWAAAWLL
						R*AVGHPEALV/DMIRWPCPIQ
						WE/YPVFC*AFLPSLLGPSFPFPR
						NNIRLTWVL/YHESSMGIFFSIG
		İ				SIHFIGQHGDVPWLHSSFYRHG
		l				KAYRFLFGFSAVSIMYLVLVLA
						VQVHAW\SLYYSKKLLDSWVH
						QHTGEEA
19224	49592	A	19336	65	665	GVRGFWAGTMASRAGPRAAG
						TDGSDFQHRERVAMHYQMSVT
	ŀ					LKYEIKKLIYVHLVIWAAAWLL
						R*AVGHPEALV/DMIRWPCPIQ
						WE/YPVFC*AFLPSLLGPSFPFPR
						NNIRLTWVL/YHESSMGIFFSIG
						SIHFIGQHGDVPWLHSSFYRHG
						KAYRFLFGFSAVSIMYLVLVLA
						VQVHAW\SLYYSKKLLDSWVH
		L				QHTGEEA

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19225	49593	Α	19337	71	1265	RCSKSLRMSTFFSDTAWICLAV
1						PTVLCRTVFCTYKKSSGQLWS
						WMVCLAGLCAVCLLILSPFWG
1						LILFSÉSCFLMHTYLSGQELLPV
						DQKAVLPTGGDCGLDHALSKY
						LDELGFTVFAGVLNENGPGAEE
						LRRTCSPRLSVLQMDITKPVQIK
						D\AYSKVAAMLHDRGLWAAIN
						NAGVLGFP\TDGELLLMTDYQR
						CMVVNFFGTVEVTKTFLPLLR*
}						SKGRLVNVTSMGGGAPMERLA
						SYGSSKAAVTMFSSVMRLELSK
						WGIKVASIQP/GEGFLTNIAGTS
						DKW\EKL\EKDILGPPPPLRVQE
						DYGQDYILAQRNFLLLINSLAS
						KDFSPVLR\DI\QQAILAKSPFAY
						YTPGKGAYLWICLAHYLPIGIY
						DYFAKRHFGQDKPMPRALRMP NYKKKAT
19226	49594	A	19338	1	861	NIKKAI
19227	49595	A	19339	248	930	DVHQLGNMAVIHSHLWEGLQE
						KFLKGEPKVLGVVQILTALMSL
						SMGITMMCMASNTYGK*PLFP
		1				CISGYTIWGSVMFIISGSLSIAAG
					•	I*N\TKGLVRGSLGMNITSSVLA
						ASGILINTFSLAFY\YSITPYCNY
						YGQPPN*LSMGTMSHLKGVW
				,		MGMVLLLSVLEFCIAVSLSAFG
						M*SALLVPPGGGCVNSAITFFT
						WAETASSTTLNEGLRGTQRFNR
						QMLPEIYA
19228	49596	A	19340	167	1164	RNLRLPSSALAGVAPSHREPRV
						RTISQTDRRKDA/PDPARRDRVE
						LAPAYTSPPRCSANVGTSTRRL
						RALCPARVRAPLRLPPPASSPLG
						SEEQRRKVPAAARMRPRAGAG
						RSTLPWLPAT/WPGLPSLPLYL*
						ICGGNNSY*R*RKGEQFE*RNSS
						*DYRYLPSSPASPNV\RERQPNE
						NGPDDKEKDRPERPSPLDTSPL
						LDPPAVSDIAGKMSDEFSLADA LPEHSPAKTSASYPDHMVLPTD
						PAAAGPLGPWGSMSSGPWVPG
						MGGQYPTPNMPYPSPGPYPVPP
						PPQASGAAPPVPWGTVPLGAW
						GPPAPYPAPTGSYPTPGLYPTPS
						NPF
L	L	LJ				171.4

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19229	49597	Α	19341	2	716	SDPSPLLDPTAVSDIAGKMSDE
						FSLADALPEHSPAKTSAVSNTK
						PGQPPQGWPGSNPWNNPSAPSS
						VPSGLPPSATPSTVPFGPAPTGM
						YPSVPPTGPPPGPPAPFPPSGPSC
		1				PPPGGPYPAPTVPGPGPTGPYPT
						PNMPFPELPRPYGAPTDPAAAG
						PLGPWGSMSSGPWAPGMGGQ
						YPTPNMPYPSPGPYPAPPPPQAP
						GAST/ILFHGAPFPPGAWGPPAP
						YPAPTGSYPTPGTLS*FPPGAW
10000	10500	<u></u>	10242	5.4	170	GPPAPYPAPTGSYPTPGTLS
19230 19231	49598 49599	B A	19342 19343	3	178 249	RHKGSPSPHQTQEPSWLHPVDP
19231	49399	A	19343	ادا	249	ALGPQVELPASLVLCARTPQPL
	1	l				GGR*DWV/PGTPAGPQESRAAP
						VPASAPPSTPRGRGGRI
19232	49600	A	19344	2	504	WAPVRPEPPRRAPPPAPQRPVP
17232	147000	l'`	17311	-		STTQGLRSAGA/PAPGLAGSYP
						CSPGYVNAPIDSLYLATFMGIR
:						VKRPPNR/PLCEQ*SFYHLGAGE
						*PGKSRVPIPQFQSP\GDPTVRV
						V\PRR*SVPPTSCATPRSWCCSD
		1				T\PGHPLGSPEQLPQPGPAAELP
		1				GGVVPRSTPPCPAL
19233	49601	A	19345	3	335	
19234	49602	Α	19346	2	447	CACTPQPLGGRWDWAPWSRG
						RRSSGRLRLHRNPRRRGKAQA
						WRAAVPRPAPREGS*GPGRPAA
						PSAGPPSPRPPGTPAGPQAPRAA
						PVPA\PRLPPHLPAS*GSGLRPSP
				ĺ		AQKGAPTVQ/PVG*RAPQVPPK
						WEPRQRRRRERARAVRTA
19235	49603	A	19347	108	729	QHAGSPQSPRLLSAPPLPGLPL
						WWHLRSPSAPPLHCGSPFLGWP
						RLEPTPSACREVWRERHKWEP
]						G/PACGACGPAGVPGGRG/PWW
						APHSEHPASPAGPG/PMGNLAP
		-				GPVAAEGVLSPPAVPAHRRCAR
						FLAGP*L/PFPRGRAWDLQPAM
						PEPPT\PPWAPV/PAEPP*RAPPP
		1				APQRPVPS\PPKGAQDWQAAPP
		1	<u> </u>	l		AAPVRDPLGEASWAPE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19236	49604	A	19348	11	816	MWESLTLSRDFLNGFDQNADS
					}	GIDNKVQVEVASDEDEELVGN
						WNKERENPFSGEKLKPAAEICIS
						YKEPNANHQDNGENVARACQ
		1				RPLQHPLPSQAQRPRRKQRFPG
						PGSGPTWDLVPCIPGAP/RLD*K
						GPRKPKSWYSEGQRYESPSFGIP
						LGSDLF\PTNALLLKPYKLPPSR
						PRDYCGRAAFPRGRVRDLQPA
						MPEPPT\PPWAPVRPEPPR*APPP
						AP/PAPSPIDHPRAEECEHTAQD
						WQAAPPAAPVRDPLGEASWAP
		<u> </u>				ESGGDVESLYV
19237	49605	В	19349	4	669	
19238	49606	В	19350	1	930	
19239	49607	A	19351	170	444	
19240	49608	В	19352	11600	1122	NOT BUILD COUNCIL BEGINNERS OF
19241	49609	Α	19353	1628	3591	KGLRLKLCSFVQLRFSIRNFRLG
ı			! }			ILTYWKKEFEIVGWPRLEPTPL
ı			:			ACREVWRERREREPGLRAALA
ı		1				GQLEFRVGVGLAGPSLGAAGQ
ı						PCWPRVIRDLAPGPVAAEGVLG
ı		1			Ì	PPAVPAHRRCAGFLTEP*LPSRG
ı						AGLGTCSPPCLSLPPTPWAPVRP
ı		1			}	KPPQRAPPPAP/PAPSPIDHPRAE
ı						ECERRAHDWQTAPPAALVPDP LGEASWAPESGISDEYITPMFSF
ı	ŀ					YKSIGELKMTQEEYALLTAIVIL
ı						SPAWATERHPVFKKRERKKPSN
ı						CLEGVIKAAAELAVGLPWNLG
ı						KFHTGDGTSAGLQGIAPNSVGP
ı						GQPAPEWMWGWERLLTDQDK
ı		}				TQGSAFSCTATFQDAQKQLLQ
ı		1				ARFIHLSGLWEQQDGLWLSNW
ı						LSLKGQIGGPALTEPLRLSVNR
ı						GTWPPKLRDARRRSAGRADRR
ı						APTGRVLPGSPGAGPQTTPPPY
ı						DWSLRNRRRPAATDPEAQAAA
ı						AAAAVGQGRVGPAQIEPSCRPA
ı						GRSDPPAPGPEAEVCWVSRLRE
ı						PVRSAFAWLRQSRALIGSWGR
İ						QGDVAPAFTCPPNAERGATGV
ı		1				RGGWGCWCGRAQLGRRDPAA
İ						TRPALNPERYQSPKSMQGLERP
ı		1			:	HVRNWGIELETLRDAFRPQPSV
İ						WRTARGEAGKRKGRLVECLGS
ı		1				
ļ.			Į.	B .	1	QRKVLEGVLLYAVTRGSFPNLF

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		
19243	49611	A	19355	951	1453	FQTQEPSWLHLVDPTPGLQVEL PASPVPCAHIPQPLGGRWDWVP
						WSRGWCSSGRLGPHRSPWSGW
						EAQAWQAAGPEPCPMGRQVRP
	-					GEKSSAVPVGQPG/SGHSVHPP
ŀ	1		1			QPLARVLSPPLPGASRAGRLLR
		}				VWGPPSPRPPGTPAGPQARHTA
						PVPARASPSTPPCKLR
19244	49612	Α	19356	111	658	LPSRGAGLGTCSPPCLSLPPTPW
						APVRPEPPQGAPPPAP/PAPSPID
						HPRAEECERTARDWQAAPPAA
	1					LVRDPLGEASWAPESGGEPTPS
						GPGVAGSGSTHTHALTHACTH
						AHTRAEPLRLPPPPRPLPPPLSPP
		1				/PPAPLLPSKPFSLSTDRISSKSR
						QQKVPFGPRDPEPRGVRAGSW
						NRTTH
19245	49613	В	19357	1	552	
19246	49614	Α	19358	1299	1813	AHFPRAGTGAACSACWPAGVP
						GGRGLGGPRTRSSRPCWPRAK
						GDLAPGPAAAEGVLGPAAVPA
		1				HRRCARFLAGP*LPYHGAGLGT
ŀ						CSPPCLSLPPTP/SGSCAPEPPRR
						APPPAPRHPVPSTTQGPPWETC
						LLSIYGNAMNDERAWSLEGFW
10047	10.61.5	ļ	10250	380	905	VLKPSDSWMTSLSERTISFF
19247	49615	Α	19359	380	1905	WIPHWGCRWSCLPVLRRALAF
						LSPWVVDGTGRRGVGSGAGRG
						GSGRTGAHGVGGRLRHGGLQV PSPAPREGS*GPARNRAQ\PGGP
			,			ALLGDSVHPPQPLAQVLSPP\FP
						GPAGLAGCSECRGPPSPRPPGTP
İ						AGPQAPHAAPVPARASPSTPPL
						VHLTNDVLRVKVKWAFFFTLH
19248	49616	A	19360	1	1052	MGLAERCPLNHLTLTVSSVKCS
17210	1,5010		17300	1	1032	FLLAHILSVAGACYGVLRGAPH
1		1				TFPGITCPGCSQENNPIKKWAE
		l				DLNRHLIQEDTQMANKHLKDA
	ĺ					PHRYIIRELOVKPTMRYCHAPT
						GVAHIQSTNTPNAGEDVEHQEL
				}		TFIADGSPVTTRGHMDLVQTSL
						SRTLGVRSTGSWSACAEFSHGY
						CLAKRLLQLLKHGNGNSNSGG
						NAECITRHCQKPLTGILSFILAL
						NKVKNTRPFDRCHSEGSEKEPS
						FCVTCPRRPEVNQPSPYMPQPG
						KSSTHLSVVQLRALEVGEPQAF
						PPVE/PEGPAGLRHPLWL*V*GQ
	1					GSGP/SSPPCLSLPPTPWAPVWP
		1				EPPRRAPPPAPQRPVP\RPPKG
L		ــــــــــــــــــــــــــــــــــــــ	l		<u> </u>	

SEQ ID NO:	of peptide	Met hod	SEQ ID NO: in USSN	location of first	codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19249	49617	Α	19361	417	1092	KWCSKRTAFSFSLQGGVEGEA
						RAGTRAACGACGPAGVAGRRG
}						LGGPPHSGQPASPAGPG/PVRDL
						APGPVAAEGVLGPPAVP/APPA
						LCSIS/PPGCSCLPAG/PRLRTCSP
						PCLSLPPPPWAPVRPKPPR*ATP
						PAPRRPVPST/NPRAEECGRTAR
						DWQAAPPAAPVRDPLGEASWA
	1					PESGGDVENLYVQLRDCKYTN
						QHPVSSSGFVNAPISTLYLVNL
						VGTWRIFMSS
19250	49618	В	19362	817	1596	
19251	49619	A	19363	350	644	CDACEDDADE A CODED CD DACED
19252	49620	A	19364	60	1362	SRVEPRVRSASQDSRSRDNGPD
						G\MEPEGVIESNWNEIADS\FDD
						MNLSESL\LRGIYAYGFEKPSAI QQRAILPCIKGYDVIA\QPQSGT\
						GKTATFAISILQ\QIELDLK\ATQ
1						GLGS*APT/RENLAQQIQ\KVVM
İ					}	ALG\DYMG\AS\CHACIRGAPTC
						VAEVQKL\QMEAPHIIVGYPLG
						RVF\DMLYR\RYL\SPKYIKMFV
						LDEA\DEM\LSRGFKGQIYGHIQ
]		ŀ				KAQAAPPRVVLLSATMPFD\VL
		1				EVTKKFMRGPH/IRILVRKEELT
						L\EGIRQ\FYI\NVEREEW\RLDTL
						CDLYETPDPSPQASHLPSTTRR\
		1				KVDWPHPRRMHALRFPWESAQ
						HEDMAQKERR/DVIMKEFRLAS
						SRV\LITT\DLLA\RGIDV\QQVSL
						SHQLNY/SFPTTRENY\IHR\IGR
						GGRFG\RKGVPINMLTE/EKTKR
						NLEDIETF\YNTSI\EEMPLNVA
19253	49621	Α	19365	911	1278	NCGIVISKHLVFTLA*MMCHFL
						ASLQRKGILKRSASV*AHSH/DP
						SFHMKIYNAGW\GHRT*HPAGG
						FAGPPPWGSVWGLAARTRENA
						PRPALPLPRPAPSGLAHGADGD
	į					SNTEKIQSRDNRKG
19254	49622	Α	19366	26	743	ETSADMAPSVPAAE\PEYPKGIR
						AVLLGPPGAGKGTQAPRLAEN
						FCVCHLATGDMLRAMVASGSE
						LGKKLKATMDAGKLVSDEMV
						VELIEKNLETPLCKNGFLLDGFP
						RTVRHAEMLDDLM\EKRKEKL\
						DSVDLNFSIPDSLLIRRITGRLIH
						PKSGRSYHEEFNPPKEPMKDDI
						TGEPLIRRSDDNEKALKIRLQA
						YHTQTTPLIEYYRKRGIHSAIDA
						SQTPDVVFASILAAFSKATS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				<u> </u>		
19255	49623	A	19367	40	1002	VASGTRTELQDRRFRDTSGDM
		1				FPSVPP*EPEYTKGIRTVLLWPP
						GAGKGTQ/SGETEV*MACSLGY
						IASNLQSLHLNDSKAPALSAAIY
						CPRHPNWAENFCVCHLATGDM
						WR\AMVASG/SSELGKKLEGNL
						WDGWGNWVE*WKWVSWSSF
		l				GEGILENPLVAKNGFLL\DGFPS
						GPVKARQK\SPMDLMGEGGKE
			1			KLDFC*FEFSIP\DSLL\IRRNHKE
						GLIPPPKKWAGSYPRREFQPLP
		l				KEPP*KMNITGGNPLIRRSR**M
						KRPLKIRPASPTHTQT\TPLIE\Y
		Ì		1		YRKRGIHSTI\DASQTPDVVFASI
1				1		LAAFSKATCKDLVMFI
19256	49624	Α	19368	793	1152	SGRKRRGCGGQSVKQAPALTP
į						GPPAPVFPIPEAPAACAPHHPGQ
						GNRRKCNFPPVCPT\PKTASSCP/
						EPLPPPPSLQR*PCPCHPWHHLE
						FSFLDEVTFSFNKTFPKRLVSSK
						MLPLKVGH
19257	49625	A	19369	1	873	
19258	49626	В	19371	1	322	
19259	49627	В	19372	64	2250	
19260	49628	Α	19373	20	368	CEARGGSGHGGRAERAGVCGG
						PAELLPKGSKEEQRDYVFYLAV
						GNYRLKEYEKALKYVRGLLQT
	1	1				EPQNNQAKELERLIDKAMKKD
		į				GLVGM/AIVGGM/ALGVAGLA
L						GLIGLAVSKSKS
19261	49629	Α	19374	3	492	
19262	49630	Α	19375	62	570	GSRGQWPSEAVLNELVSVEDL
						LKFEKKFQSEKAAGSVSKSTQF
	į					EYAWCL\VRSKYNDDI\RKGIVL
					,	L\EELL\PKGSK\EE\QRGFTSFYL
						ARGGGTTRLQGITEKALKLRSA
				,		GLLAAQEPQEQPRPKETGNGLI
						DKA/L*KKDGLRGAWPIVGGM
		1				ALGVLAGTCRSSSDFAVFQ

SEQ ID		1	SEQ ID NO:	1		Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	Sequence		027510,217	sequence	or pepme ouque	positive nucleonal insertion)
19263	49631	Α	19376	1	944	MACHLQAGSPGNLVVYFQSKP
			•			EGLRTREPMMSVPVQVKGPST
						RKLMMSEDPLTLGNWHELRVS
						RTAKNGILQVDKQKIVEGMAE
						LQQVVGKNMDSGVGPVVHGG
						GLTRTGYLLNPESLQQKAFERR
İ		}				QIQGCDEKLACVGKQTLGTWP
	İ	İ				LSESPSLTRLYCRRQLTQGSLR
						MSLNNAGVPGQTAPDQRSGEG
						VAAPAGPGEEVGREEGEGAAA
						GKRQCPCQQGSLCVDTSRPRLP
						KSQAQIPSSSRSSPSSSEQQRGQ
						LPSVREHVEGGQGEERAPHCD
	ļ					GGGRVLLASGLAWLGTENIMD
				1		LASSPKDDWGVAVPSRGCERPS
						TFTLVLVPSPIVRRRACSGQPISP
						GGGQETHGADTSHLTNP*ICLL
						SKAFCCRDSGLSK*PVLVRPPPC
£						\PQALPQSPCSYLLLAVALPFPP
	<u> </u>	_				LSSAYPPVRFHSLLCEIHEARAS
19264	49632	Α	19377	1177	1603	WVSSFSSSSFSSFPPPPPPPPPPPP
						PPSFLPPPPPPPP/PPI*DRARRCQQ
	İ					TSIPLNKLNINKLLLLLLSPRPD
						DREKFTPQPQSARQREPIKGTA
						SRRGEGPICPTLSLYFLHVFLCF
						SCLSSQIHIYILRLSTVLVLIKNS
19265	49633	A	19378	1	896	FPFSLLPPPKMPKGKKAKGKKV
						APAPAF\VRKQDAQE*WLNPL\F
						KKRP*DFR\IGQ\DIQPQKRPPPL
		l				L*KWPR\YIRLQR\QRAILYK\RL
						K\VPSAITPVSPRALDPANQLLS
			ŀ			LLKL\AHKYRPRDKAQRSKQR\
						LLA\RAEKKAA\GKGDRPNQRD
						PPVLRARS*HPSPTLVEEQEKLS
						WVVIA\HDVD\SIELGLSFLA\AL
						CRKNGGSPYCIIKGKGQDLGR\
						LVHRKDLAPLVAFTQ\VN\SED
						KRRFWLKLVGSYQGPNYNDRY
						RYDEIRRHWGGNVLGPKSVARI
						AKLEKAKAK\ELATKLG
19266	49634	A	19379	3	491	AKEALDWEEGHQ*DQRL*SRN
						SQLGICGGLRGFRGDRAASVFS
						R*AEKESVRQL*ARVPGALGVA
						ARANKHQRRWREGDSTQTATN
						TVDRRATRGRPMEWRATRVAD
						EEDGDRRRGDNRPQARTFRPRT
						SVRRAAGTNNSEGRRPTNAGT
				L		NGGRGVDIYTVN

SEQ ID NO:	SEQ ID NO: of peptide	1	SEQ ID NO: in USSN	Nucleotide location of first		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19267	49635	A	19380	1	1501	MRRSTPGVLPAAVGGARVRRA
						VGLSGTGAEAGRAGAMVEKEE
						AGGGISEEEAAQYDRQIRLWGL
					İ	EAQKRCHRMKEALSTAPFTDK
	ļ					ETHPKVSLAQVTRSSASVMVY
						GAFEVIGOROSSAAKPRRSOSE
						SLGPEFQGLWEWLPASQHIMRS
						LKKQDGMGQGYRVSLEWEVG
						QLIFLEEKVTP\EDPGAQFLIRT\
						GSVGRNRA\EAS\LERAQNFNP
						MV/DMLKVDTEDIEKKPESFFT
						QFDAVCLTCCSRDVIVKI\DQIC
						HKNSIKFFT\GDVFGYHGYTFA
						NLGEHEFVEEKTKVAKVSQGV
						EDGPDTKRAKLDSSETTMVKN
						KVF\FCPVKE\ALDGWILDACDE
						MQMAALKRTTSDYILLQVLLK
						F/RVTDKGRDPSS\DTYE\EDSEL
						LLQI\GNDVLDSLGISPDLLPEDF
						VRYCF\SEMAPVCAVVGGILAQ
						EIVKALSSAGTLP/PTTNFFFFR
						WA*KGEWGLWECLGPQVNSG
						FGQPQRCQLQHAHLYSLFPFFM
19268	49636	Α	19381	2	411	PGTSQGVFSPHRLGGAVRSPCP
						PPQDSRQASAGGAESQ\LKQRG
						FPGE*GCRSPVSCAWPQNRMG*
						GPQHREAGAWSGE*KTPIWLK
						QERCNRPGLLVTAELAAPAPRP
						DGMAPGCGGIRCTPGRPRRLLH
						SGPPSP
19269	49637	Α	19382	165	309	
19270	49638	Α	19383	1	825	LATQPEGAGRVPSIHPQGGPEA
						ALQDLRTLMGSSRLHSGKESIG
						HNSSAGSSSQGPHHPRRKTAIL
						EKGGKQPHRMLPFSPGLGSSPE
						PANAGSPDTWLCPP/AAARVL*
						NPVKPRKGRSEPRSGWASQLPG
1	ļ			-		GDSRLPLRPGTSQGVFSP/APAG
						*AGKLVLGVLSLSLKQRGFPGE
						*GCRSPVSCAWPQNRMGCRTSP
						GPSPTSPTAAGE*GSRPRRRKPG
						RVQGLQRDGPTLTQEDALTAA
						KRCPLPVTSVPQTPETSTGAAA
						PGSQPVGAPPSR
19271	49639	A	19384	1	378	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19272	49640	A	19385	381	1653	QRPRSGTFTMGRKSLYLLIVGIL
						IAYYIYTPLPDNVEEPWRMMWI
						NAHLKTIQN\LATFVELLGLHHF
						MDSFKVVGSF\DEVPPTSDENV
i						TVTETK\FNNILV\RVYVPK\RKS
						*STKEGGLFYIPCGGWRRGSYS
						LIVYDLPSSWTVE\RPGSSLVSN
						HL\RLAT*F\HFPLQLGELYDAL
						R*FLRKKVLAKYGCEP*GESGIS
						G\DSAGRGI*LSAV\TQQLL\DDP
						\DVQDPNFKIQS*IYPALSASWM
						*VLPSYQGKFKFSYFYSKSLQ\C
						RFWSEYFTTDRSLE\KAMLSRQ\
						HVP\VESSHLFKFINWSS\LLP*E
			:			GLLKGH\VYNN\PNYGSS\ELAK
						KYPG\FLDVRAAPLLADDNKLR
						GLP\LTYVITCQYDLLRDDGLM
						YVTRLRNTGVSG*LNN\HVEG\
						GFHGAF\SFLGLKISHRLINQYIE
19273	49641	Α	19386	3	667	DAWADAWVPVSSEVAYLFGG
						CHTCGGLPVAHPMMPRGSRSR
						TSRMAP\PASRAP\QMRAAPRPA
						QVG\QPPAAAPPSAVG\SSAAAP
						RQP\GLMAQMATTAAGVAVGS
						AVGHTLGHAIT\GAFSGRK*LPE
						PARP*HHFNQEASG/TPPAQHSK
						QQRHPHLQQPCP\YEIQTVFWE
						FCPGTRGDIQAL/CEGFNE\VLK
]					Q\CRL\ANGFGPNEEVSTFGRD
10074	10640	<u> </u>	10207	0.1	1057	WKFSSHNLS
19274	49642	Α	19387	81	1257	RHQSDQTVPRLSRPHVQSQKSS
						MAAAAKPNNLSLVVHGPGDLR
						LENYPIPEPGPNE\VLLRMHSVG
			:			ICGSDVH\YWEYLS/RFGNFIVK
						KPMVLGHEASGTV\EKVGSSVK
						AP*NQGDRVAIEPGCSPEEN*W
						NSC\KMG\RYNLSPSIFFCATPPD DGNLCRFYKHNAAFCYKLPDN
						VTFEEGALIEPLSVGIHACRRGG
						VTLG\HKVLVCGSWANRGWVT LLVAKAMGAAQVVVTDLSATR
						` '
						LSKAKEIGADLVLQISKESPQEI ARKVEGLLGCKPEVTIECTGAE
1	}					ASI\Q\AGIYATRSGGTLVLVGL GSEMTTVPLL\HAAIREV\DIKG
						VFRY\CNTWASGGFSMLCVPKS
						V\NVK\PLVTHRVSSWRKVLEA
19275	49643	Α	19388	2	431	FETFKKGLGLKIMLKCDPSDQN
17413	47043	Α	17300	2	1,21	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19276	49644	Α	19389	2	326	ARGSTRLMESRVQHL*CTRKEL
						TPEVLQEWLDELEEMMLVVH
		l				MPRFRIEDGFSLKEQLQDMGLV
			į			DLFSPEKSKLPGIVAEGRDDLY
		1				VSDAFHKAFLEVNEEGSEAAAS
19277	49645	Α	19390	2	368	
19278	49646	Α	19391	1	2123	MAQFSIVGMCLGLFNQSPTDGS
		1				GVQLFVIMNQVMAANLAETHA
		1				ALCPTLSSGTSARFRGKNQFSG
						GLPQITLSPLAQPCGRLAAMYS
			,			NVIGTVTSGKSLPSTSPLVPLLL
				}		SGFAPLFLKGLQNWTQTTQEG
		1				KSPSAAPASSQPQACPTDHVRE
		•				SALGACLSLPGEDGKIQKVYLL
,						SLLLIGFWDCVTCHGSPVDICT
						AKPRDIPMNPMCIYRSPEKKAT
]	EDEGSEQKIPEATNRRVWELSK
į						ANSRFATTFYQHLADSKNDND
						NIFLSPLSISTAFAMTKLGACND
						TLQQLMEVFKFDTISEKTSDQIH
						FFFAKLNCRLYRKANKSSKLVS
						ANRLFGDKSLTFNETYQDISEL :
		Į				VYGAKLQPLDFKENAEQSRAAI
	1	1				NKWVSNKTEGRITDVIPSEAINE
	1			ļ		LTVLVLVNTIYFKVLRMALERP
						QGLPLALQLTPFFFKWRDRSPE
	1			ļ		RANGLPKATQGLWKSKFSPEN
						TRKELFYKADGESCSASMMYQ
						EGKFRYRRVAEGTQVLELPFKG
		1		ŀ		DDITMVLILPKPEKSLAKVEKE
		1			į	LTPEVLQEWLDELEEMMLVVH
						MPRFRI\EDGFS\LKEQLQDMGL
]				VDLFSPEKSKLPGIVAEGRDDL
		1				YVSDAFHKAFLEVNEEGSEAA
						ASTAVVIAGRSLNPNRVTFKAN
						RPFLVFIREVPLNTIIFMGRVAN
						PCVNLSEALAVLVVNLTTLMK
						RTHGESLDFMERKRRPAESILL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19279	49647	Α	19392	9	1630	PALCPTLSSGTSARFRGKNQFS
1,72,7		l'`	1,,,,,,			GGLPQITLSPLAQPCGRLAAMY
						SNVIGTVTSGKRKVYLLSLLLD
						SFGDCVTCHG\SPVDICTAKPRD
						IPMNPM\CIYRSPEKKATEDEGS
				ł		EQKIPEATNNRRVW\ELSKANF
						PLLLTTFLFSTWADSKD*H*LTF
						FCSPLSIFQGFLLWTKV\GACND
						TLQQLMEVFKF\DTIS\EKTF*SR
						SHFFFA\KLN\CRL\YRKANKSSK
						LVSANRLFGDKSLTFNETYQ\DI
						SELVYGAKLQAPGTSRENAEQ\
						SR\AAINKW\VSNKTEGRIHRCS
						FPSGRPFNELTVLGGGFNTIYFQ
						GACWK\SKFSPENTRKELFYKA
						DG\ESC\SASMDVTREGKFRYSG
						AWLEGT/QVLVVCPFKGDDIT\
						MVLILPKP*EGAWAKVEK\ELT
						PEVLAKSGWD*FWREMMLVV
						HMPRF\RI\EDGLQV*REQPAKN
	İ					MGPLSD\LFSP\RKSPQTPPGIVA
	:					EG\RDDPPMSSDAIPIRHFLEVN
						EEGSEAACKYPLL*LPGRS\LNP
						NRVTFQGQQAFPWVFIREVPL\
10000	12612	<u> </u>	10000		1505	NTIYLHGAELANPCV
19280	49648	A	19393	3	737	
19281	49649	A	19394	1	705	
19282	49650	A	19395	2	757	
19283	49651	A	19396	1	1764	MANUAL I I TO DECOMA OG
19284	49652	Α	19397	1	710	MAWALLLLTLLTQDTGSWAQS
						ALTQPASVSGSPGQSITISCTGT
						NNDVGSYNLVSWYQQHPGKA
						PKIMIYEVSKRPSGVSNRFSGSK
						SGNTASLTISGLQAEDEADYYC
						CSYTSSRPD\VVFGGGTKLTVL
						GQPKAAPSVTLFPPSSEELQAN
						KATLVCLISDFYPGAVTVAWK
						ADSSPVKAGVETTTPSKQSNNK
						YAASSYLSLTPEQWKSHRSYSC
						QVTHEGSTVEKTVAPAECS
19285	49653	A	19398	1	679	
19286	49654	Α	19399	1	743	
19287	49655	Α	19400	2	736	
19288	49656	Α	19401	2	718	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19289	49657	Α	19402	1	711	MAWALLFLTLLTQGTGSWAQS
						ALTQPASVSGSPGQSITISCTGSS
						SDVGGYNYVSWYQQHPGKAP
						KLIHYEGSKRPSRVYDRFSGSM
						SANTASLTISGLQADDEADYYC
						CSFAGSSALRIFGSGTRVIVIGQ
						PKANPTVTLFPPSSEELQANKA
						TLVCLISDFYPGAVTVAWKAD
		ŀ				GSPVKAG\VETTKPSKQSNNKY
						AASSYLSLTPEQWKSHKSYSCQ
						VTHEGSTVEKTVAPTECS
19290	49658	Α	19403	3	720	
19291	49659	В	19404	1	711	
19292	49660	Α	19405	105	858	RRQDSGQSPAWPAGALLLTLLT
						HCAVPGSW\AQSVLTQPPSASG
						TPGQRVTISCSGSSSNIGINDVT
						WYQQVPGTAPKLLMYSDNHRP
1						SGVPDRFSGSKSG\TSASLGPSV
[GLQSEDEADYYCGTWGLAA*M
	ŀ	İ				VFMSSAGGTTLTVTSQPKAAPS
						VTLFPPSSEELQANKATLVCLIS
						DFYPGAVTVAWKADSSPVKAG
						\VETTTPSKQSNNKYAASSYLSL
						TPEQW\KSHRSYSCQVTHEGST
						VEKTVAPTECS
19293	49661	В	19406	63	756	
19294	49662	Α	19407	1	757	MAWTPLLLLFLSHCTGSLSQAV
						LTQPSSLSASPGTSASLTCTLRS
						DINVGSPFIYWYQQKPGSPPQF
						LLRYKSDSDNQQGSGVPSRFSG
						SKDASANAGILLISGLQAEDEA
						DYYCSSYVGTNNFGVLFGG\GT
		ł				\KLTRP*GQPKACPPRVTL/FPHP
						PSEEASKPNKA\TLVCL\ISDFLP
						GKP*QLPWKAA*QPPSLAGVET
		ŀ				PPHPPTRSNNKYAASSYL\SLTP
						EQWKSHRSYSC\QVTHEGSTVE
		l				KTVAPTECS
19295	49663	Α	19408	1	224	YISLPAGVLSLACCTLYGISWQF
						DPCCKYQVEYDAYKL*RLPLH
1			ļ			TLTCSRPTRDNRVFRSPHSVRQ
		1				QSLGRCLA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last Amino acid sequence (X=Unknown,			
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide		
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)		
				sequence				
19296	49664	A	19409	235	955	ESTVQVSVFCHRVPARTSSQTA		
19290	49004	^	13403	233	933	LPVLLGLLGFTVSGPLNLPEPSV		
						IVGNGELVGPQTCLHCGMEIYN		
						G\ENAQ\DQF*VTELGAGP*EGP		
		ŀ				VTSYIVIE\PTRIGDETARWITVG		
						NCLHKTAVLAGTALLLHPGLR		
	•					CTLRFIPHHISPAPLGVL\SLACC		
						NL\YGIS\WQ\FEPFACK*QVEVR		
						TPNKLSRLPLHTLTLLPPRVVLV		
				ļ		RKDDLH\RKRL\HNPIGTGPPLV		
						YCVK\KIYELYAVMIFK		
19297	49665	A	19410	1	764	MRLGIGFACPWKGLADEFDLKI		
19297	49003	^	13410]'	704	EKQRNQGRLLGFKLQHLVPFTE		
						MEKMGGDRQKHRTAPRQAER		
						RRSGRVSFQPPLQPLDVTLLPG		
İ						ATREGNCPRASRGRRPSPISHCQ		
		1				ETQGPEGRSDCTIRGVLSSIPAA		
						RPLRKGSGFPTVCEKRPEQGQQ		
						PALGP/AGPAAVPAPAASRTQW		
						GSRCPCGAGARRGYWRGLLGL		
						CNLISPLDPGSPGSGRWGSAGA		
						LSRPADSCPPWWPRAGA*RRG		
						AAAEAGNSPSRCGASRPAEEQS		
1						GAFAGPLPSSPSQ*RAPGAEV		
19298	49666	A	19411	209	536	TRSSCVTLVC*LAMVSLRVARP		
19298	17000		17411	209	330	HGVPGTMFGKRKKRVEISAPSN		
			İ			F/EHRVHTGFDQHEQKFTGLPR		
						QWQSLIEE/SARRPKPLVDPACI		
						TSIQPGAPKEVCSPRASGTRSR		
19299	49667	A	19412	124	1983	1510. 511115 5511615 11616		
19300	49668	A	19413	424	1582	PPWAAVHGDRPHRVPGTMFGK		
13300		1.	1,,,,,			RK/KRVEISAPSNFEHRVH/TGF		
						DQHEQKFTGLPRQW/QSLIEES		
			1			ARRPKPLVDP/ACITSIQPGAPK		
		i				WGETGSWPPF/RHLPEG*HGPPI		
						PGCQREPQRVSHEQ/FRAALQL		
		İ				VVDPGDPRS/YLDNFIKIGEGST		
		j				GIVCTPPSGKLVAVKKM/DLRK		
			1			QQRRELLFNERPWQCP*HP/RR*		
		İ				EAQVPLPVLQALSV/LHAQGVI		
						HRDIKSDSIL/LTHD/GQVSKEVP		
						RRKSLVGTPY/WMAPELISRLP		
						YGPEVSPGWLGCPAVDSVRCH		
						FQLQRWREMGPSLGSHCQNGA		
1						LDKEEGQRPSHTGSLQWGSLG		
						VERTLRVNMYECVHACNSFYL		
						VNKLFQKDSTQEASVTMVPDP		
						DKDGKNRASQKRKAHSQGARR		
L	<u> </u>		1		<u> </u>	DI DOMINIO QUINTINO CONTINU		

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19301	49669	A	19414	488	707	TCGEGLSWTRPLPTASSLQV\H* SLSAGERPYLCDYPDCGKAFVQ SGQLKTHQRLHTGEKPFVCSEN GVNLVF
19302	49670	A	19415	16	454	GPAGRCPADCSEEKEGPSA/GD SGRPPSCT*PWKGLPVMAILPG EAPVPGPGWPEPDAGG\PLWR WGSPSPCAAALHRP\SGTTPPGR ALSGPGLERAKQRSPRVSPAPR AAPHGHPTHCLPIPAGPRPAKD HPAYGPPASCHHHLAAS
19303	49671	A	19416	166	1191	TAGSLVLAARGT*GPPPRARK* GPAGRCPADCSEEKEGPSAETP GTTPPGRALSGPGLERAKQRSP RVSPAPRAAPHGHPTHCLPIPA GPRPAKDHP\PTHQKPPPRTEGS TGGHCCGPPIFLIPNTSMLTVYP SGPVMPAGFTLSPSDTRPSLLN GQEVLRKMNGAVQYLSLCDYII PTCQCLATTVLCSVSMSLTVFA VSYMVELLLLQNAQVHQLVLQ NWMLKALPPALQDPPHVPPRV PRAARTRLPAV

2421

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19304	49672	Α	19417	1	2481	MRSGDLKFGKSAAWPSGQKEK
						LVFRGKIEEGLEYLHKKKPSAN
		i				SQDNGKEASKAFHGPLQQPLLS
						QALGPDREEWFPGQAPWPHCC
						VHPQDTAGCIPVVPAPAMAQR
						CTGTAWVTASEVAPSLDCFHIV
						LSQQVYRIHRPTSSLHPQYGKA
						TRHTTPAVSMRAAAGAEPCKA
		ļ				TGADWPKTLGAQPSHPCTMDV
						GQGFKKDDSGAALAPMAALM
		İ				GWAGIECLWLFHTEGVNCWP
		ļ				WDLGEKNGLLGQPRGPAAVCS
	į					LRTLLPASQQPQLLLRPWLKDA
						QVQLGSLLQRVQAIGLGGFYIV
:	İ	1				LSQFIDPSKASYPPWVKNSRHS
		1				TAAQPMRAAAGAEHCKATGA
		1				ELPKALGAQPSCPCVLDVGQGF
		1				KKNDFGAVGLNNWPAGFWNV
		ŀ				MGTCKSRLCFVLLSGKKSSFRV
		1				EVLTQCLDSHTLETYQFYNREV
	į.					AFEDVLESSQTGLVVQKPQTDR
						LTKRGRGEATRNLRGLVPSMP
						RSLLPSTAQPTAWKCQRGPSPY
		-				THQDMALIPSPTARCLSPAKEP
	1					KQEEVGEKSLLPDPTLPLTDPR
						LTGSTEQAHAEGLAALMSALR
						VSHLQGRGGVVTLVDSQLGVI
	1					AVSSTQFNKGPSYRLLADVQN
1						RLLPKYDSQKEAELRSWIKGFT
						GLSIRPDFQKGLKDGIILCTLVN
						KLQPGSVPKIN\ASV*NWH*LEN
•						LSNFLKAMVSYGM/NPVDLFEA

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19305	49673	A	19418	1	1299	MHQEDLRAWYLDLGLPSHQNE
1						TTDVSVVYERFSLTDATYPKQA
						KQWDFPFLRVSTAQPTAWKCQ
						RAPSPYTHQDTALIPSPTARWL
						SPEKEPKQGEVGEKSLLPDPTLP
						LTDPRLTGSTEQKKDLAAGPVG
1						SALRRRPPAVSSTQFNKGPSYR
						LLADVQNRLLPKYDSQKEAEL
						RSWIKGFTGLSIRPDFQKGLKD
		ļ				GIILCTLVNKLQPGSVPKIN\ASV
						*NWH*LENLSNFLKAMVSYGM
						/NPVDLFEANDLFESGNNMQVR
		l				VSLLALAGKAKTKGLRSGVDIR
						DKYSKKQNFDDTTMKASQCVI
						RLQITNKCASQSGMTAYGTRR
1						HLYDPKNRILPPMDNSTISLQM
1						GTNKYASQVGMTAPGTQRHIY
		l				DTNLGIDKCENSSMSLKMGYT
						QVANHSGQVFGLGRQIYEPKY
						QPGGPVAHGAPSAGNCPGPGE
19306	49674	Α	19419	135	290	HREEESKSSVFLNPRGVQGTGV
						GDHRMILGRGRAVLSKE*AAG
		<u> </u>				PRGGRHES
19307	49675	Α	19420	92	1058	KGQYLNPNRGKGGPEPPFEPPV
						PPASRSDMAQNLKDLAGRLPA
						GPRGMGTALK\LLLGAGAVAY
						GVRESVFNVE\GGHRAI\FFNRI
						GG\VQQDT\IL\AEGFHF\RIP\WF
						QYPH/IIYDIRAKTSKKSPPLQGS
		ŀ				KDLQM\VNISPAKCWSRPNAQE
		ļ				LS*ACTSRLGLGLTRER/RCLPSI
		ļ				VNEVLK/AVWVAKFNCLTA*FT
						QRAPGYSLLDSAGSLTERAKGL
						SSLIL\DDVGQSQKLSFSPRSYT
						A\AVEAK\QVAQQEAQR\AQFL\ VEKAKQ\EQRQKI\LQGRG*GR
						GLPKMLGEA\LSKNP\GYIKTSQ
		1				RFEPAPEIILQRRIAHIHRNP
19308	49676	A	19421	3	861	LDDSWEPVHPGCGDFICQWSEE
19306	49070	^	19421]	801	SGLWSTAEVWRERRGREPGLC
		l				EGARGTARVPGGRGPPHQSGQ
		1				PVPPAPGS*GA*\PGPAAVEGVP
						GPPAVPPASAALEFSRGFSCLPR
		l				CHLVVTKIGKQEDDVSIKRKNR
						PDYWPAPEPVAGCEGSDQEQV
						KPNATLEKHFLTEGTGPRSPSC
						LSWPPSGPHAAADPAMVPETPE
						PDRPQLTKKFCEARVMAVGTS
						NVLGQTLKPSRTGGTSWSWVF
				[YLLTANRLPFDVKRKGGGPSSI
						KPRPHYDHRLLQDFKEQVIHHF
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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19309	49677	Α	19422	808	923	
19310	49678	Α	19423	1	339	
19311	49679	Α	19424	1	621	
19312	49680	Α	19425	3	1836	MAAVEAAAEPVTVVAAVGPK
						AKDEEEEEEPLPPCEA/LRWAP
				į		VLWRGPAWATAFLEEATAEEP
						GAAPGSPPDSPDRTLRRLRAER
						RRLDS/SLLALSSDFAQVQFRLR
				:		QVVRGAPAEQQRLLRELEDFA
						LKG/IPDILGYEGPGDPASDEGD
						GLPGDRPRLRGEDQSEQEKQER
						LETQREKQKELILQLKTQLDDL
ľ						ETFAYQEGSYDSLPQSVVLERQ
		l				RVIIDELIKKLDMNLNEDISSLS
		1				TEELRQRVDAAVAQIVNPARV
						KEQLVEQLKTQIRDLEMFINFIQ
						DEVGSPLQTGGGHCECKAGGK
						TGNGCSRTGSSRTPPGNSRTKA
						EDVTKVRKTGLHLMRRALAVL
						QIFAV\TRLVVPQARSLQPCGRG
						SRLTETTLPC*R/ELEASVDRVK
						QLALTAEPHDHVITSANLQDLS
						L*GKDELTMAVRKELTVAVRD
						LLAHGLYASSPGMSLGMATIA*
	1					LLPAFSSVPEAMHPRELFVKYY
						HAKNGRAYVESPARKLSQTFA
						LPVTGGTVVTPKQSLLTAIHMV
						LTEHDPFKRSADSELKALVCM
						ALNEHRLVCWMNLICKSGSLIE
						PHYQPWSYMAHTGFESALNLL
						SRLSSLKFSLPVDLAVRQLKNIK
19313	49681	Α	19426	366	477	
19314	49682	Α	19427	3	432	WNSRDAGYEFDICFTSVHLY\AI
						RTLWTVLDAIDQMWLPVVRT
						WRLNERHYGGLTGLNKAETAA
						KHG\EAQVKIWRRSYDVPPPPM
						EPDHPFYSNISKDRRYADLTED
						QLPSCESLKDTIARALPFWNEEI
						VPQIKEGK\RVLIA

SEQ ID NO:	of peptide	Met hod	SEQ ID NO: in USSN	location of first	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	or peptide sequence	deletion, \=possible nucleotide insertion)
19315	49683	Α	19428	1	890	MRTRRTPRGEHSLSAFSSCVDK
						ISRVCKSFGLELENSIQTLWAER
						EQSICLVAAGHLVSSSGMRNW
				Ì		CHARRHAAYKQVLIRHGEVAW
				ŀ		NLENRFGGWYDANLNPAGDEE
						AKRWEQALPDAGYEFDIC/FKR
		ł				TIRTLWTVPDAIDQMWLPV\NV
						SKDRRYADLTEDQLPSCNSLKD
						TLARALPFWNEEIVPQIKEGKW
						SDHCLGALVIPVKDGCNLHGD
						VTTVSSLTPEERALEEPLNRPTN
						A/P*HQEG*CWESHQKPVPFQT
		l				LGCLCEFSRGASRRKQARVVFR
						ERSNLCRSISGGEGF
19316	49684	Α	19429	1	443	RALPFWNEEIIP\QIKEGKRVLIA
						\AHGNSSGAIVKHLEGLSEEAIM
						VLNLPTGIPIVYELDKNLKPIKP
						MQFLGDEETVRKAMEAVAAL
					_	VCVLESLPALFHGSEMPLGPVQ
		İ	:			VYLSLISESCSSCLTLPHGSSVH
						LSITVLNPFSISV
19317	49685	Α	19430	24	1039	AVLALLRGRRGGLLLRNLLIPV
						GAASPARRQWPPYKLVLG\RHG
				1		ESAWNLENRFSGWYDADLSPA
						GHEEAKRGGQALRDAGYEFDI
						CFTSVHLY\AIRTLWTVLEMPL\
						SEMWLPVV\RTWRLNERHYWG
						S*PGLN*KQKLAAKHGLRPRLK
						I/WRRSY/D/VPPPP/MEPDHPF/Y
ł						SNISKDRRYADLTE\DQLPSC\ES
						LKDTIARALPFWNE\EIVPQIKE
						GK\RVLIAAHGNSLQGHCSRLL
						EGLF\EEAIHGSLNLPDLVFPLSI
						ELDKNFESLSSPMQFLG\DEETV RKAMEAVACPGQGPRSEGRRG
						RILSPGCTLPCPVFVPSSPSHLA
						HVTLDHICKTS
19318	49686	Α	19431	86	431	VYVSTMSLSCPASELAVRRQR
15510	77000	^	17431		731	WSSQFQDHPPGPLQPRPV*EML
						LL*PTLDLQPVVPPAVAILSWKP
						SALTISSQIVTPGLKTRSFFTRKT
						FMKIWNHRQKYQKTMLVMFP
				·		RYPSL
19319	49687	A	19432	599	900	ATWLFISPTTTTCQLSKTLLVPL
		l .	.5.2			NFLEKQYLTFLLFLEFPTFSLFF
	}					RHTEDHLMRSGCLKVCSISPLS/
	1	ŀ				PLPPSSSHVRHLTPSLPFAMIVS
						FLRAPQKLSR
19320	49688	A -	19433	210		HFRKIFTVFQLKLVEYNLLSGH
1,7520		l' `	17.00	[-··	[STQQSHPGGTVLL*DLVCPVV
		Ц		L	L	OT SACTOR AND DEACHA

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19321	49689	A	19434	239	464	SPKGGPPNPSHGTKFHTRPCEE TTKQALCEEHGCLFHLGAGGLS PKRESAKGDKGGASVHVVCHI S*SPCS*LGP
19322	49690	В	19435	1	838	
19323	49691	C	19436	80	443	
19324	49692	Α	19437	1	775	
19325	49693	Α	19438	209	469	
19326	49694	A	19439	304	375	NVGE*SERHPCYD*TPREVCPPS P
19327	49695	Α	19440	879	1188	
19328	49696	Α	19441	615	917	ATWLFISPTTTTCQLSKTLLVPL NFLEKQYLTFLLFLEFPTFSLFF RHTEDHLMRSGCLKVCSISPLS/ PLPPSSSHVRHLTPSLPFAMIVS FLRAPQKLSR
19329	49697	A	19442	674	1554	GQMASSSVSLQQGVPQAADPG WVTAGASLTRAWAAGVQGDG PGTESSISASSRRVLRLGMTRTL LWACVLLWVCILLLGLCPAVG PVS\SVGPVSYCGPVPYCGPYPA VGPCPAVGPVPCCGPYPS/VGP VPCCGPVSCCGPVSCCGAVSCC GPVS\SVGPVSC/VLGLCPAVGP VS\S*CGP*PAVGLCPAVGPVPC CGPYPS/VGPVPCCGPVSCCGPV SCCWACVLLLGLCPAVGPVS\S VGPVPCCGPVPCCGPVSCCGPV S\VVGPVSCCGPVSCCGPV CDHSSHSLLLTFTML
19330	49698	Α	19443	4688	4824	

2426

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	l .	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19331	49699	A	19444	1	2274	MLKPLMGGGAHRRAAAGARK
19331	43033	^	17444	I	2274	WDGTAIIASTSGLEVLWVTTSR
1						SKTVESVGLAKPGQHNDARMN
						_
						LAIALTAARYGAATANYMEVV
					1	SLLKKTDPQTGKVRVSGARCK
						DVLTGQEFDVRAKCVINATGPF
						TDSVRKMDDKDAAAICQPSAG
						VHIVMPGYYSPESMGLLDPATS
						DGRVIFFLPWQKMTIAGTTDTP
		j				TDVTHHPIPSEEDINFILNEVRN
ŀ						YLSCDVEVRRGDVLAAWSGIR
						PLVTDPKSADTQSISRNHVVDIS
						ESGLITIAGGKWTTYRSMAEDT
						INAAVKTHNLKAGPSRTVGLFL
						QGGKDWSPTLYIRLVQDYGLE
						SEVAQHLAATYGDKAFEVAKM
						ASVTGKRWPIVGVRLVSEFPYI
						EAEVKYGIKEYAC/DCCGYDFT
]					ļ	SYSPGLS/EMSRQQRKPYPGLL
						N*/CGRELNWDDYKKQEQLET
						ARKFLYYEMGYKSRSEQLTDR
			:			SEISLLPSDIDRYKRRCHKFDAD
			ŀ			QRGFMTVVEVQRVLESINVQM
			•			DENTLHEILNEVDLNKNGQVEL
			:			NEFLQLMSAIQKGRVSGSRLAI
						LMKTAEENLDRRVPIPVDRSCG
						GLYFCVRNWWVLGLTDFKNE
						AADPSGVKLQTFTVSVTARKGS
						VDPKNSGAQLASPSGSRTGAA
						GGAACQSRALRLHSSALGWSM
						GLGAVEHGAVLIGEARAAQEP
						MEGVGGSGMAGCRSQVLPRG
19332	49700	В	19445	439	1542	
19333	49701	Α	19446	2	418	CLSLPPTPWAPVRPEPPRR/CAT
						PCSTAPSPIDHPRAEECERTVQD
						WRAAPPAAPVRDPLETGKIDQE
						IHKYNTPGFTGCLSRVQFNQIAP
						LKAALRQTNASAHVHIQGELV
						ESNCGASPLTLSPMSSATDPWH
						LDHLDS
19334	49702	A	19447	111	215	GTCWELE*RKGKPIFWGEIKAS
					·-	CRNLHKLQGAKC
19335	49703	Α	19448	740	959	DLYPHHLSWQSRNAARKRKS*
1,7555	1.7703	'		1, 10		RGQRPIPAIS*GR\SQKQQQNVN
						TSTPSHCPRLPYECRPLLHGAQS
						HRPPKG
19336	49704	В	19449	260	508	TAG TAG
19330	I	Г	17777	1200	200	<u> </u>

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19337	49705	A	19450	11	490	RLSTSPPDSGAQLASRSGSHTG AAGGAACQSRTMRSHSSALGW SMGLGAPLQRH*INSSGGSWLV DKESFPVPSALNFSPLGRKSSPC SKVLYMPNPVSHSMIPSGSERS MDTNGSCWYLTCSLSAPGRRD TRIRKVFSQGRLLVLTLMISPSV GHSTA
19338	49706	A	19451	42	263	SGVSPMSFNTFTSAPAFNRSWT TCLCPK*HARCRGVQPKLDLLL PLQLMSAITSFLAMVSNSCLTSS ALPFLA
19339	49707	Α	19452	662	1564	GSRICSFQTRSGHPECPQCHWT RQNHSSIQQPHKAAYTKKTTIH PPNRSNSCSSIWCKLQKLYMCG P*GRSFNFSANLHQPQKGHSGE RPF\EKNKGRAILLKSHRVHTQE MPFPCSKVGKGYLVSSGLFQH QAIHNEKPCRSAMYGDMFHTQ QGHFKCIDYGEAFSPKDTPGQH QIIHTGEKPYVCTECEKTCTRSS NLIQHKPKGLLSLLVLSPSTGN ALFSTLSQLTQLLLTFDVRSDA TFFRKGSGRHQLWLALPEDAR ARVTSVSGTRRRDHGGARGVV CLIHGCVPDTRSSPDVW
19340	49708	A	19453		1167	MIREQKVKEVYICAGKKKSHM HLWHNGEKYNRLGRKDRKMT HEGWLIKIPHFPGRKYKASGLH WEQMKHMGGRGKVHLEFTEM YVAGRRGEVKDEVLEPMSLTE VYIPTENDREVWRERRLRAPGL CAALEGQLEFRVGVGLAGSAL EAAGQPCWPRAGRARDLQPAM PEPPPASVGSCAARASLRSAAP CSTAPSPIGHPRAEECERTAQD WQAAPPAAPVRDPLGEASWAP ESVMWPPLDSRQSEEVSVSKRH MDATDKIKFPWLKEKRTIEFGE AAGYVCHRRYKDMEPHEKMG EAAKKVHWVQMRSRARDLQP AMPEPPTHSKGSCAARASPT/T HHPLLHDAQSHRPPKGLRSAGT WRGTGRQLHLQPQCGIHWVKP AGLLSLL
19341	49709	В	19454	118	559	

SEQ ID		1	SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence	1	09/540,417	sequence	of peptide sequence	detection, \-possible nucleotide insertion)
19342	49710	A	19455	26	813	LSHYLACSLRLPSGSRIRSAWLI
						PSIMLLNYTSPGLPRHRSKSENR
						LDDTARRSPGMAGPGCPPLPGG
						VPDVRAAYGAPTSAQRPRVGW
						PPGGRWPRGARGARSFLKKPFR
		1				GTPTQVRGAWQCPPPPLTV\EK
						*PWHPCGRFQGRRHEEKGGRE
						LGHSKSPPPRRSHYPAGEKGEE
						ERKEREESSCAGLRRGAHEHA
						QGQSLPGELPAPGDPHPGRCVG
		İ				GQGRSRMGPCS*SLAQHPPVCT
						RGQREKQERRATNKPVRGAPK
19343	49711	A	19456	497	756	TWLKSGISFPPALFTIAKTWR*P
						KCPSMIDWIKKMW/HIYTMEY
						YAAIEKDEFMYFAGTWMTLDT
				ļ		VILSKLSQEQKTKHRMFSLISGS
19344	49712	Α	19457	100	819	EWSSVRRILVEKRALRRPHPQC
					İ	LCFRMKTILSNQTVDHSQKNV
	-					DITLKGTPQ*SCKGPRGT\LR\RD
						FN\HIKCGTSALLGKEQQRGFR
						VD\KWWGNRKGNWATRSGLF
		İ				GSHVQDHDPRGVTTGASGYKD
						EGPVYASPSPHPTVVIPGRNGSS
						LLKSRNFLG\EKYIPQGFRMKTR
						VLLCSVSQSPEKMKLILE\GNDI\
İ						ELVFKFQRALIQQ\ATTVKNKG\ IRKFFGWVSMSLEKGTVQPGLI
19345	49713	С	19458	395	574	IRRITOW VSMSLEROT VQFGLI
19345	49713	A	19459	1	1476	
19347	49715	A	19460	2	323	
19347	49716	A	19461	539	711	HRPFPQDRKAGECLLHEYEDLV
17540	13710	``	17101		,	PIRDTLRLFPGGRYLPRAKHVA
1						PSEPDPGWLVLG*FYC*FCHSP
ł						ALLSGFCLAVNVYRERSV
19349	49717	A	19462	66	2873	SARTISYDYYQNWGRDGGPRSS
1.75.17						GGGYGGGPAGGHGGNRGSGR
						GGGGGGGGRG/WQGPASRAPE
						RPRNRHVVREKTGAEEQGSWR
						GKREL/LVHMDERREEQIVQVL
1						DSGQAENG*RVRTTDF\RFAPE
						DHGYGTEVSTKNTPCSENKL/G
						HPGKEVDKSRKKNV/SESGTDH
						ILTRDSEYLLQENEPDGTLDQK
						LLEDLQKKKNDLRYIEMQELV
						NLIDNHQVTVISGETGCGKTT\Q
						VTQFILDNYIERGKGSA
	1		l	<u> </u>		VIQFILDNYIEKGKGSA

SEQ ID NO:	SEQ ID NO: of peptide	Met	SEQ ID NO:	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19350	49718	Α	19463	1	2112	FRKTTRRLRKVKGRTAGCARF
						RKWLLRQHGLRTLGLICNHTSR
						MTRCRWSAESDSGDEEEEGPIV
						LGRRQKALGKNRSADFNPDFV
						FTEKEGTYDGSWALADVMSPL
						MKKRAATTLHEKIEKARKRRK
						TEDKEAKSGKLEKEKEAKEGSE
:						PKEQEDLQENDEEGSEDEASET
						DYSSADENILTKADTLKVKDRR
						EEEEERTGSRRIF*RCIS/ATMKT
						SSFQDMNLSRPLLKAITAMGFK
						QPTPIQKACIPVGLLGKDICA\C AAPGTGKTPPFALPVLERLIYIP
						SPGSSL/TRVLVLVPTRELGMQ
						VHSVTRQLAQFCNITTCLAVGG
						LDVKSQEAALRAAPDILIATPG
						RLIDHLHNCPSFHLSSIEVLILDE
						ADRMLDEYFEEQMKEIRMCSH
						HRQTMLFSATMTDEVKDLASV
						SLKNPVRIFVNSNTDVAPFLRQ
						EVIRNRPNREGDREAIVAALLT
						RTFTDHVMLFTQTKKQAHRMH
						ILLGL\LGVQVGELHGNLSQTQ
		-				RLEALRRFKDEQIDILVATDVA
						ARGLDIEGVKTVINFTMPNTIK
		ł				HYVHRVGRTARAGRAGRSVSL
						VGEDERKMLKEIVKAAKAPVK
		1				ARILPQDVILKFRDKIEKME\KD
						VYAVLQLEAEKEMQQSEAQI
						NTAKRLLEKGKEAVVQEPERS
						WFQTKEERKKEKIAKALQEFDL
10251	40710	ļ. —	10464	254	116	ALRGKKKRKKFMKDAKKKGE RQHAH*ESSPLPNLTYPGTQILR
19351	49719	Α	19464	354	446	KAAGTSA
19352	49720	A	19465	42	388	RAAUISA
19353	49721	A	19466	177	497	
19354	49722	A	19467	271	773	TAMSSEEGKLFVGGLNFNTDE
	"					QALEDHFSSFGPISEVVVVKDR
		1				ETQRSRGFGFITFTNPE\HASSC
		1				Q*RSQ*TGEVSWNG\RQIPLLDH
		1				AGKSCSGEPRGRWALGPMGVG
		1				SQLILEVVGDQG\YGSGRYYDS
						RPGGYGYGYGRSRDYNGRNQG
						\GYDRYSGGNYRDNYDN
19355	49723	Α	19468	2	414	GRVGPPRRAGGPRASAAGDPA
						RGGAAPKMSQTAMSETYDFLF
						KFLVIGNAGTGKSCLLHQFIEK
						K/WSVTRSYYRGAAGALLVYDI
						TSRETYNALTNWLTDARMLAS
						QNIVIILCGNKKDLDADREVTF
	1	<u> </u>	<u> </u>	<u></u>	<u> </u>	LEASRFAQEN

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO.	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19356	49724	A	19469	397	445	GHRGAHRPRTLRSVVVRRAHR CSYSGIWD/TAGPDRFRSVTRSY
		1				H*GRPGALLVYDITARETYNAL
						TNWLTDARMLASQNIVIILCGN
						KKDLDADREVTFLEASRFAQE
						NELMFLETSALTGENVEEAFVQ
ĺ						CARKILNKIESERMGSGIQYGD
ŀ						•
19357	49725	A	19470	2	559	AALRQLRSPRRAQAPNAQECG
19357	49726	A	19470	1	483	
19359	49727	A	19471	95	352	RKATKMATIWASGTPKQPCPR
17337	49121	^	19472		332	ELLAQERRQTRTRYNQ*DQ*QE
1						RRSRQRIMGVGCLHMLRSEKA
			ŀ			PQQRSFAHCTPKSSRPSARLT
19360	49728	В	19473	247	1364	I QUELLINGSIG SMELT
19361	49729	A	19474	422	877	CHTQTQ*QHTP*QPTPIDMAPTS
	' ' ' ' '					HARMPFHTHFLTQILTLVSRSP
						DPG/PHGGKVPTRIPIPPAQPWG
		İ				PSTLPSSSFLGPPRPFPSPPVPGP
						GGENRKPFTSPP*A*\GFPGPPSG
ļ						P*AGGEG/PPVVGRGATPSPPTP
						SPACGPLKCWWHLIKN
19362	49730	A	19475	168	522	LALHPTRPLIPATDDEV\TVTED
	1,3,23					KINAPH*KQPVVNVEPF\APGLF
						AKA\LANVNIGSLICNVGAGGP
						CCQPAGGCTQPGGSCPLHCWL
						LPA*GRRKWEAKKEGFRGSFD*
						LTWGFGSF
19363	49731	Α	19476	1	328	
19364	49732	Α	19477	1	1204	MASRLLLNNGAKMPILGLGTW
						KGSILRVGFGSSSREPPPYRGQP
						LMGGPPVRILGPSGRPKRHNAR
						GRRGKWASQTGGPRAQTGTWS
						RRQGQVTEA\VKVAIDVGYRHI
			1			DCAHVYQNENEVGVAIQEKLR
						EQVVKREE\LFIVSKLWCTYHE
			1			ERGLV\KG\ACQEDTSSDLKL\D
						LPGPLTLIHWPTGFKPGKEFFPL
						DESGNVGPSDTHISDTWA\AME
			1			KLVDEGLVKAIGISNFNHLQVE
			1			MILNKP\GLKYKP\AVNQI\ECHP
	1		1			YLTQEKLIQYCQSKGIVVTAYS
						PLGLS*TGPWPKP\EDPS\LLEDP
						RIQGGSQPKHNK\TTA\QVLIRG
						SPCKRNLVVIP\KSVTPERI\AEN
						FK\VF\DFELS\SQGYDPP*LSLPT
						RNWR\VWCPCWSCT\SH\KDYP
					-	LPMKEVLKA
19365	49733	С	19478	300	489	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19366	49734	A	19479	243	471	CDHLQVSRPRIPATVPVEWKRS EMDSKEEEKEKKEKKEKRRR *RRRGKEEEEERRRRKRRKK RGRKK*IKKRV
19367	49735	A	19480]	365	MGLYKDLLDTRYLFYLVPGYG LTQIELEVIAKDSAKPVGTPRV GLKLEMGIYLSGHLDKFIFAMP FKIKDILSLKKEEEK/ERRRRR RRKEEEDEEEEEEEEERRR RKRRKKKRKKKID*KKEEEKE EEEGEEGEKKKKMKKKRKKK KKKKGGGGGKGGERRRGRKK
19368	49736	A	19481	96	508	PAGSLVCPNQCSSARLFSSFFCS SAVMSASPAVGKRGWAELTGS FLPKPPLKTLGSGREDR*GQVW WLTPVIPALWEAEVGGSLEGQE FETSLGNMVKPLLYWLGSNGW LWKLGRRLWQLLWLQDDWPT RGSHILVL
19369	49737	A	19482	3	1013	SRPSWGTHWNEYARRLTLAPS WPLP*GHPVSPSLRVSCL/PHYY ESVLEKKGPCDRDYKKFWAGL QGLTIYFYNSNRDFQHVEKLNL GAFEKLTDEIPWGSSRDPGTHF SLILRNQEIKFKVETLECREMW KGFILTVVELRVPTDLTL\LPGH LYMMSEVLAKEEARRALETPS CFLKVSRLEAQLLLE\RYPECGN LLLRPSGDGARQRVGHHAADA QRDA/PWFRHYKVEAGRGPKY VIRCGTAVLLHLPWTTVVNYF VSHTKKALVPFLLDEDYEKVL GYVEADKENGENVWVAPSAPG PGPSPLHKVPPSRCQPASSQDKL PPLPPLRNPGREIT

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	i	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19370	49738	Α	19483	63	1496	LISAMAC\ARPL\ISVY\SEKGESS
ŀ			:			GKNVTLP\AVFKAPIRPDIV\NFG
						SHPNLRKNNRQPVCWQGN*AG
						HQT\SAE\SWGYWPRAVGSKFP
						R\VRGGGTHRS\GQGCFLETMC
						R\GGRNVLHPTKTLGRRWH\RR
	:			ļ		SSTQPKKRIRPSGSAPGCPQPLP
						ALGHGLKGHLY*GKFPETSFW*
						VER*KLKG*QEEPREAVFAPLR
	•					KLKRPGNDIQKGSMASSAK*RA
						GQKAKLRKPVARIPAAGAPCII\
			:			YNEDNGYSFKAF\RKHPWGLL
		İ				MLNC*GKLEHF*KLAPLVGHV
						G\RFLHLGLESALREV*DDLY\G
						T\WR*SHFPLQGVNLPIFSPWHK
						\MI\NTDS*AESLKSPEIPREPLRA
						PR\KKIHSQSS*RRNPLEKL*EIM
			ļ			LKA*TPFAKAHAAGTPFFGQAR
						N\HKLRVDKAAAAAA/ANLQA
						NSDEKAAVAGKKPVVGKKGK
			:			KAAVGVKKQKKPLVGKKAAA
						TKKPAPEKKPAEKKP\TTEEKKP
						AA
19371	49739	Α	19484	3	730	KRLPKMAEVQVLVL\DGRG\HL
			<u> </u>			LG\RLA\AIVAK\QVLLGRK\VV
	:			1		VV\RCEGIN\ISANF\YRN\RLKY
						LGFSPPSGMNTNPSRRPLTTSG
						APSRIF\WRT\VRGMLPPQNPSG
						GQGRLLDRLQGV*RASPPPYGQ
			•			EKSGMVVP\AALKVVR\LKPTR
			1			KFCLIWGRL\APEVG\WKYQAV
						TAPPGGERGKRKAKIHYRKKK
	:		}			QLMRL\RKQAREETWRKKIDK
						YTEVLKTHGLPGLSPIKTVNSS
	1					KKKKKKRAAA
19372	49740	Α	19485	1	660	PTRPRTRGRRTRKSTPMKGLGA
						RYREYKV\VGRCLPTPQNATRG
						PLLPACEIFAP\N\HVVAKVPAF
						W*LCILS*KKMKKFFQGKIVYL
						WGKVF*EGSPLRVERTFGDLGL
						RLLTPRERAPHNIVTGEY\RGP*
		ĺ				PTAGGCSTQLLTRDNGVARHR
						RPEAHFHFRFMEGWEGDRGP/C
						KIPPGGLVKASSNDSKIKVSRW
						PHRV\LAAFRNKARFTT\KRPNT
L	<u> </u>		<u> </u>	L	L	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19373	49741	A	19486	51	471	EAGRLPGNILLDSIMACGLVAS NLNLKPGECLRVRGEVAPDAK/ S*VLPARCVHGILSDRLVGQWG WSWPRWPHAHPLPPSFVLNLG KDSNNLCLHFNPRFNAHGDAN TIVCNSKDGGAWGTEQREAVF PFOPGSVAEVCI
19374	49742	Α	19487	59	527	ETQSWPCGLVA\SN\LNFNTLES CLRFPKPSWLPDA\KSFRC*NLG QRPPTTWCLHFQPSASTAH\GS ARHASLCNSQGTAGHEGGPTQ REAVFPFQ\PGKCCRGVAFTFRT RANLNRSSLPDGITEFQVSPNGF NL\EAINYMAA\DGGLSRSKCV AFD
19375	49743	Α	19488	2	440	KQQCRDTGDEVVCSCFVGYQL LSDGVSCEDVNECITGSHSRAF GESCINTVGSFRCQ/RISSCATG YELTEDNSCK\DTLGSFRCRPKL QCKSGFIQDALGNCIDINECLSI SAP\CPIGVT\CINTEGSYKCQKN VP\NCGPLVNHLKE
19376	49744	A _	19489	2	2061	
19377	49745	Α	19490	2	2120	
19378	49746	В	19491	63	255	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19379	49747	A	19492	2	1941	PPPMERAAPSRRVPLPLLLLGG
						LALLAAGVDADVLLEACCADG
ļ						HRMATHQKDCSLPYATESKEC
						RMVQEQCCHSQLEELHCATGIS
				1		LANEQDRCATPHGDNASL\EAT
						FVKRCCHCCLLGRAAQAQGQS
						CEYSLMVGYQCGQVFQACCVK
						SQ\ETGDLDVGGLQETDKIIEVE
						EEQ\EDPYLNDRCRGGGPCKQS\
						CRDTGDEVVCSCFVGYQLLSD
		1				GVSCEDVNECITGSHSCRLG\ES
						\CINTVG\SF\RCQRNSSCGTGYE
				i		LTEDNSCKDI\DECESGIHNCLP
İ						DFICQNTLGSFRCRPKLQCKSGF
						ITRML*GNLLLDIQWSVLSIQLP
						RCPIGD*HGI\NTEGFYNVPEGT
						CPNCGPLGYHLQRGRGTRCV\D
						V\DECAPPCLSPVGKG\HR\CVE
				•		LFPGSFPLAECKDGVTIFDGISR
						DVLFDVQRSCQRLPRGACC\GH
						K/CTENTLGSYL\CSCSVGF\RLS
						VDGRSCEDINECSSSPCKPGSVA
		ł				NV\YGS\YQCY\CR\RG*PASAIV
				1		GWEFTCERTFDGVAALPHRGH
		1				TFWFLTAGIQTSWGSFPVAACP
						SSGLTGWAPNGPATCQ\DI\DEL
						C*LGIHNCFHSTEDLVSNIPGAG
						FRCLGLSSCPENYRRSAAT*S*G
						NSCMRAIGAGWEKEKGKLAGS
19380	49748	В	19493	123	237	
19381	49749	Α	19494	285	456	VINKRLVCTRLLRRKWSCLQPH
		1		1		RMM*VQAGSRGSQPKYHRHE
				<u> </u>		MGWTGHTDTPPHRL
19382	49750	A	19495	383	693	NFLPAQPCLSAPPS*S/RSLLDNS
		1				PLNCKAFHSSSHHLLKLESCSE
		1]			NSPGPLL\QPGGTCLGACPGCFS
		1				FSPGISAGKPLSKAASPPVCTGS
						GAETSTTPPSPLR
19383	49751	В	19496	28	236	

SEQ ID NO:	SEQ ID NO: of peptide sequence)	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19384	49752	A	19497		927	PAAPQVAGRLGLGCPLHLHVF AVVSAMLPLLRCVPRVLGSSV AGLRAAAPASPFRQLLQPAPRL CTRPFGLLSVRAGSERRPGLLRP RGPCACGSGCGSLHTDGDKAF VDFLSDEIKEERKIQKHKTLLK\ MSGGWELELNGTEAKLVRKVA GEKIPVPF\NINNSIPP\TFDGEEE TLARARRFEDREPELTSLPKF\V VE\VIKND*WQERPLVLD\VHYP EDEVGQEDEAESDIFSI\REVSFQ S\TGESEWKDTNYTLNPDSFWT GPLYD\HLMGFP*PNRGVDNTF
19385	49753	A	19498	3	342	A\DE\LVELSTAPRSPPGVPLPFP LDTNLPANRVPAGLEKRLCAA AASILGKPADRVNVTVRPGLA MALSGSTEPCAQLSISSIGVVGT ARDNRRH\GAPFFGFSPRELALG QDRILIRFFPLESWQNGKIRTGM TFL
19386	49754	Α	19499	3	520	SSARAVSVPLPAMPFLELDTNL PANRSARGGWRKRLCAAAAS\I LGKPA\DRVNVDRYAGGFAIGA *AGST\EPCA\QLS\ISSIRRQWGT AED\NRSHSAHFF*VSSPRELSP GARDRIILFRFFPLESWADWAR *GTVIDFFMIWGTEGFQGHLVN LAASFQKRSSWAEVKGLG
19387	49755	Α	19500	7	355	-
19388	49756	В	19501	172	479	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	1 ··· • · · · · · · · · · · · · · · · ·
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19389	49757	Α	19502	1	1656	LEDPDSSSRPRRAGSGHSASFPF
				•		PPLEQLWRQVKTLNPKAEVAR
						AQAALAVNISAARGLQDVLRT
						NLGPKGTMKMLVSGAGDIKLT
	}					K\DGNVL\LHRNCKFQHPTASL\I
						AKVSNKPQE*YNLVDGTDFLIV
						LINWESLLKQA\DLY\ISEGLHPR
						IIT\EGF\EGAKEKALQFLEEVKV
						SRE\MDRETL\IDVARTSLRTKV
						HAELADVLTEAVVDSILAIKKQ
						DEP\IDLFMI/SVIMEMKHKSETD
						TSLIRGLVLDHGARHPDMKKR
ŀ						W\RDAYILTCNVSLEYEKTEVN
						SGFF\YKSAEEREKLVKAERKFI
						EDRVK\KIIELKRKVCGDSDKGF
						VVINQKGIDPFSLDALSKEGIVA
						LRRAKRRNMERLTLACGWG/M
		İ				PLNSFDDLSPDC\LGHAGLVYE
						YTLGE\EKFTFIEKCN\NPRSVTL
		:	:			LIKGPNKAHTSLRIQRWQ*GDG
						L*GAV\KNAIDDGCVVPG\AGA
						VGSGQWAGSPGFKHKPSC*RA
						RAQLGSPKHLLDAFAHYFPKVL
						CFRNSGFLTFQGNIKLKFKAE\H
						SESG\QLVGVDP*TPGEPNWWP
						ARSKAVWG
19390	49758	Α	19503	2	424	EGPVMAPPNPGPATLGGPGPDP
						DLGRLPLHEVFLHIRVPARPRG
						APLHRRGLRGRHAVRAVRQRR
						REPEDGAAGAVDRAGGAGVLG
						RGDRESEGPLTD*PREPADRAP
						LLQPERGRFSHPLPTLTCAPSPII
						FPVAERWG
19391	49759	Α	19504	3	174	
19392	49760	Α	19505	2	153	LPLHEVFLHIRVPARPRGAPLH
						RRGLRGRHAVH*PSEPGDPARL
						LQPERG
19393	49761	A	19506	3	354	EDGRHGAPNPPPATLGGPGPDP
1		ĺ				DLGGLPLHEVFLHIRVPARPRG
						APLHRRGLRGRHAVRAVRQRR
						REPKNGAAGAVDRAGGAGVL
						GPGDTEYEGPLTD*PSEPGDPA
						REPKNGAAGAVDRAGGAG

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19394	49762	A	19507	3	1246	GSTHASARMAVMAPRTLPPAT LGGPGPDPDLGGLPLHEGF\LHI RVPARPRGAPLHRRGLRGRHA VRAV\RQRRREPEDGAAGARG* EARKGPE*LGRG\NTESEGPTHR LDPSEPGDPARLLQP\E*GRFFTP FQIMYGC\DVGSDGAFSSAGYP QERLTNGKDYIALNE\DLRSWT AADMAAQITKRKWETPPMRRR SLRAYLDGTCVEW\LRRYLENG KETLQR\TDPP\KTHMTHHPIS*P *RPL*RCW\AL\AFYPCGDHT*P WQRDG\EDQTQGHGSLLETRPA GAG\TFQKWASMVVPSGQEQR\ YT\CHVQQ*RVLPKPLTLKM/W EPSSQAHPSPIRGASFAGPWFSL GSL*ITG\AVGRCW*C*RKEQLR WRRGER*KKEGAYSQGCKASD
10205	10762	ļ	10500	1	202	QCPGASDVSLHSFVKVLRQLAL
19395 19396	49763 49764	A A	19508	1	392 220	
19390	49765	A	19510	3	500	
19398	49766	В	19511	28	134	
19399	49767	A	19512		1662	MAASVVCRAATAGAQVLLRA RRSPALLRTPALRSTATFAQAL QFVPETQVSLL\DNGLRVASEQ SSQPTCTVGVWIDVGSRFET\EK NNG\AGYFLE\HLAFKGTKNRP GSALEERRWRAWGAHLNAYST REHTAYYIKALAKELPKA\VEL LGDIVQNCSLEDFTD*GRNVDV ILRE\MQEN\DAS\MRDVVCNYL HATAFQG\TPLAQALEGPSENV RKLSRADLTEYLSTH*QGPSKW CWAAAGREWEHQQLFRTFAQE GTSVGIP\WTYAEDAVPTLTPCR FTGSEI\RHRDDALPFAHVAIAV EGPGWASPDNVALQVANAI\IG H\YDCT\YGGGVHLSSPLASGA VANKLCQSFQTFSICYAETGLL GAHFVCDRMKIDDMMFVLQG QWMRLCTSATESD\VDRGKNIL RNALVSHLDGTTPVCEDIGRSL LTYGRRIPLAEWESRIAEVDAS VVREICSKYIYDQCPAVAGYGR SLGLPFHLCSRTAAQLTEQPVK APSSSPQSPGCQGVLAQKTCPSS SLTACQGAFINTLHPSGPIEQLP DYNRIRSGMFWLRF

SEQ ID			SEQ ID NO:	1		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	l .	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19400	49768	Α	19513	2	1693	AQSTLARRNRADCSWKMAASV
						VCRAATAGAQVLLRARR\SPAL
						LRTPALRSTATFAQALQF\VPET
						QVS\LLDNG\LRVA\SEQSSQPTC
						TVG\VWIDVGSRFETEKN/DSGA
						GYFLEHLAFKGTKNPALAVPW
İ						EKEVESMGAHL\NA\YSTREHT
						AYYI\KALSKDLPKAVGALGDI
		İ				VQNCSLEDSQIEKERDVILREM
						QENDASMRDVVFNYLHATAFQ
						G\TPLAQAVEGPSENVRKLSTA
						DFTRYL\STHYKAPESVLPQTEG
			ļ			VRTEQLLDL\ARNHRVHIP*TY
						AEGAL\PTITPCRFTGSEIRHP*D\
]						ALPFAH\VAI\AVEGPG\WPAPD
ŀ						N\VALQVGNA\IIGHYDCTFGG\
						GVHLS\SPLG\SGACGPTKLCQS
						FQTLS\ICYAEDGLCWVHTLFC
		1				DRMK\IDDMMFVLA/QGHWMR
						L\CTSA\TRSEV\ARGQKNIPQK
		l				MPLGISS*DGNYSLCVEDITPAA
1						LLTLWARRNPPGLKWES\RIAG
				1		GGFQVVVR*DLAPRYIYGPSCP
		}				AVA\GYWPH*SMLPDYNRIP*P
İ		1				ACFWVAFFRAGKPMVKQKGR
						GGGLWSPPPPQTQHFRLLLTCC
19401	49769	Α	19514	46	422	
19402	49770	Α	19515	568	1567	GPWAANGVVLAFPPCLGSPPAP
						HLAPQKAVPPKWGISSLCSFHF
1						HIYQVGGVAFCGRIRFLSSTDF
						KRRAGGSGSLMSQEEKHELSG
		ŀ				KIRSQIMDDHARPYLYTMSNCP
						CWAGALGPQESKCSRGRLLHR
	1	l				AFPSLV\TLLLAGQATTGLLSCY
			İ			QQQGAGWDKLDSHRPRNLAA
						WEEPCGMEAFPKPSQSLVEAKI
						AQLATPGLLDCKAACPWGAPC
		i				PRGP\MQNATKVIGNM\TEDHV
		1		}	1	DAPGSRNAEPP*KVYPP\LKGSF
						P/EKNLRHLKNTMETIEWKAFE
						NW\MHHW\LLF\ELSKQSLWEQ/
1						QSPLDAPPK/ENSLELEDPSSGL
	1.055	_	10.51	<u> </u>	1.400	GVTKQDLGPVPM
19403	49771	В	19516	1	1428	

SEQ ID NO:	SEQ ID NO: of peptide sequence	l .	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19404	49772	A	19517	1	2293	 MARSEAAAAGPGPGPPGLAAR
	ŀ					ERRGGRGCVRGCERASVRVWS
					1	VALRRARRAPARPYVARCRRV
ĺ						RRPARVAIGCGGVSSPAGVSGC
						ARVSPERPPLRLCVWALSRAHG
						GAVCRAGPGPDSLKRGLTSHPP
		ĺ				AFPLHHGSRTLLLGPLSPPPAAL
						PFLLLLWAGHLVAS/ACPGRCIC
						QNV/SPTLTMLCAKTGLLFVPP/
						SIDRRVVELRLTDNF\RRHFAN
						MTSLV/QLTLSRNTIGQVAAGA
l						F/VDLRALRALHLDSNRLAGC/
						GGDQLRGLGNLRHLI/VGNNQI
İ						RRVESAAFD/DLLSTVEDLDLS
i						YNNLEA/LPWEAVGQMVNLNT
		ŀ			ł	LT/LDHNLIDHIAEGTFVQLQSW
1						A\RAQAAHPLTVSFGGNPLHCN
		:				CE/LLWLRRLTREDDLETCASP
	ļ					N/NLTDRYFWSIPEEESCTRCQR
						FCG*ASLVAAELTSNSVLIRW/S
						TQRPVPGIRMYQVQYN/SSVDD
						SLVYRKYSLSTYMNPTR\MIPST
						SQTFLVNDLAAGRAYDLCVL\A
						VYDDGATALPATRVVGCVQFT
		l				TAGDPAPCRPLRAHFLGGTMII
		l				AIGGVIVASVLVFIVLLMIRYKV
						YGDGDSRRVKGSRSLPRVSHV
		1				CSQTNGAGTGAA/AGPGPAGPG
[PLRGAARGGVP/WLPPPSPSRPR
						PWRPR/LASAEPEVVLGRSLGG
						SATSLCLLPSEETSGEESRAAVG
						PRRSRSGALEPPTSAPPTLALVP
19405	49773	В	19518	97	255	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19406	49774	A	19519	1321	1728	EHAHDRDLHHAHYVQATLVR ASLREAVVGSPTDAWLQQRTA RSSASLLLARNRPATSVWLPRA GFIAGTGGRRERRGSEGWEGG GAGGTGAGLSAHSQTSACEGS/ CGGSAGRGRCLLSAATPPLQPP RPAPPTAGLVQPRQVPTAPPRA SAGNEHHQGWSGRL*KSLAAA RVGEAPGKLPRPRDFRKQSNGA LARASLRVSAQPGTCSSSAAAF PPLGPAPLAAPARSCDESGPRQP DGRGGPVPGQQQRGA/ESEPCA AATRPWVNRRR/TSRREARTS VACT*CAWCRSRSCACSHSPWS SASSSSAAICSSSPRA*STSS*RT GGRLRRWRRWSWGPTDPYAP AATAPTPAHFASP/ACAPHYQR LGEANLSESIISLD/CRPFGKKRP VSPSPSESPHAWQSEESAQTLN SAESGKKRAIRAQNFG/RLLLT WNAGRPTVRRPFSTRRWAKLW GQVERAGRGETQRH*SPCDRKS DLLKATQQTHGVSQIRVRVRSP LFSLLPPCPRKGGANKIPLNCKR
10407	40775	_	10520	102	447	ARFKKGFVFVCLEIQMFE
19407 19408	49775	A	19520	192	591	
19408	49776 49777	-	19521 19522	281	421	
19409	49777	A	19523	174	512	
				67		
19412	49779 49780	A	19524		1166	MGAKDGRLLSESSQGLSSCGAS LVPFCVSKSRLLMKTPVRLDKE TVSTVDSGAPTDLAQLPTVLKQ PCCSVMASGQFVNKLQEEVICP ICLDILQKPVTIDCGHNFCLKCI TQIGETSCGFFKCPLCKTSVRK NAIRFNSLLRNLVEKIQALQASE VQSKRKEATCPRHQEMFHYFC EDDGKFLCFVCRESKDHKSHN VSLIEEAAQNYQGQIQEQIQVL QQKEKETVQVKAQGVHRVDV FTDQVEHEKQRILTEFELLHQV LEEEKNFLLSRIYWLGHEGTEA GKHYVASTE/PTVERSQEAH*FP VCLLASYDEISGQGASSQDTKT FDVALSEELHAALTSAHPDLILS QDEIVTLNLASQGGSQKRGNPR RFYRRGFLEVEPLTGFVELHSS GSQCQALIESGSREDLPVCPNK
19413	49781	Α	19526	203	359	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19414	49782	A	19527	135	466	DPAGSSAASMSSAAGLCALTPP
						/GLLSLGLLLLPLVVAF\ASAEA\
ļ						EEDGDLQCLCVKTTLPGSVPGH
						IHQLGG*FKGRDPHCPTCPT**A
		l				TVEGMGRKILLGTWQAPLLQG
19415	49783	Α	19528	46	468	TGPTHASAAILVPRSLHKMPGE
						ATETVPATEQELPQPQAETGSG
		l				TESDS/GKSPASDTYIVFGEAKIE
				,		DLSQQAQLAAAEKFKVQGEAV
						SNIQENTQTPTVQEESEEEEVDE
						TGVEVKDIELVMSQANVSRAK
						AVRALKNNS
19416	49784	Α	19529	337	509	ISQQLLVVLAGINLRSWLPQKL
		1				CKYLKQDVLTALDLSVEVPEFC
1		l				PFVDHLVVALEEEGPAL*GHPA
1		1				SDTYIVFGEAKIEDLSQQAQLA
						AAEKFKVQGEAVSNIQENTQTP
		l				TVQEESEEEEVDETGVEVKDIE
		i				LVMSQANVSRAKAVRALKNNS
						NDIVNAIMELTM
19417	49785	Α	19530	91	1019	GELLALLGHRVPTRPWASGQK
ł						GVCGWCPGKSTLQGLRCGPTIR
						ACRDGSSCSVEGGRGVRLVLG
						TASQDPEKEMPGEATEPVPA\T
		1				EQGVAASPRAETGSGTESD\SD
						ESVPE\LEEQDSHPGQPHNKAQ
						AGRQQSEIDEDPVS\KAKQSRN*
						KEGPGRAMSQTGVFGQVTGVT
		1				KEFTI\RKSKNILLCHPPKPDVY
						QGAPASRYFTIVFGGSPRSEDLS
		1				PSKHNLASCLRNSKFQGEAVSN
						IQENTQ\TPTVQEES\EEEEV\DE
						T\GVEVKDIE\LVMSQANVS\RA
						KAVPSPEERTSNDIVNAIMVSV
19418	49786	Α	19531	1 .	261	MWGHFHSHPGTGQARAYTDTS
						GSPIPLIMHVNRGRHSEIPVLER
				1		NLLHFRMDAST*NLGMSSLME
						AILKNQFRHVSLAVSFAFSGEK
19419	49787	Α	19532	277	349	
19420	49788	Α	19533	792	1014	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19421	49789	A	19534	849	1752	GESRGWERRKCVRPLSSYATHL SVCDCHIGNVREGADPAGGAG *L*GARQQQTHTHKPAKATQR GEDREEKKRETKREERLGQQE NERDERGRREKEPRNRGTKRRE AEKGRRR/ESEGKREEKGEEER QERQQKRAREKRRGRGARRRG REKKKTQRPRTQAQRPPLKKIK DYTHHTTQTPTQQHHTHTTKH KQPHTNNTTVFLFFFLLAKEHR RAARKQHEPERGRGGGSKAGA KIKHTSIAAAADRRHDRPTQRD TRRKEGRRAQSGKHGRTAHKT EAGGGIPVHSTSRVQVILLLQPP
19422	49790	Α	19535	139	470	
19423	49791 49792	A	19536 19537	32	434	NRRKGA\GPVFQRAT*KH/RV*R AGLRAVDFA\ERHGYIKGIVKDI IHNPRR/GAPLAKVVFRDPYRFK KRTELFIAAEGIHTGQFVYCGK KAQLNIGNV/LP/VGTMPEGTIV CCLEEPRLCRRKRTSAEGLNKV CVYAT
19425	49793	Α	19538	10	443	
19426	49794	Α	19539	63	376	
19427	49795	A	19540	170	368	
19428	49796	A	19541	175	427	
19429	49797	Α	19542	1727	2262	
19431	49798 49799	A	19543 19544	695	596 1337	GFGTQGERRRQCRLRVTQPLG ASSATQKTPALPETPVKAEQSC TLLPSPRTPGHLQPRLLDPSPW AVLIRG\QRKGAGSVVRRATLR HGK/GAVRRLLRRWISPERHGY IKG\IVKDIIHRPRSRRAPPLRQR WFFRDS\YRF*ER\TELF\IAARG AFHTGPSFVYCRQERPSFNIGQ WCSPVGHHCLEGYNSCCLPWK EETLETCGKAGPGAFKGN
19432	49800	Ā	19545	1	813	
19433	49801	A	19546	227	381	
19434	49802	A	19547	135	740	PQPLPPPTSMARHVFLTGPPGV GKTTLIHKASEVLKSSGVPVDG FYTEEVRQGGRRIGFDVVTVVR HPGGLLSRVGLEPPP/RENVNAE LGSYVVDLTSFWSSWALTRLE GM*ADCQQWAQGQRVCVIDEE LGKMEALSVK\FSFKAVRQTLS YPRDI*SL/VHKSQVPKGKPLAL VEEIRNRKDVKVFNVTQ\ENRN HLLARYS

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19435	49803	С	19548	251	574	
19436	49804	Α	19549	2	450	
19437	49805	В	19550	721	3048	
19438	49806	A	19551	3	396	
19439	49807	A	19552	2	450	
19440	49808	A	19553	2	1152	GPPLHCTRAAGIRHEAPLGLAG LCRAADAGLCSAGWGFVKVV KNKAY\FKR\YQVKF*KTPERV KRDYYARKR\LVIQDKNKYNTP KY\RMIV\RVTNRDINLSRLLYA PV*RGDMDSLAQR*LHTELAKI LVVEGLALTNLLPAALFVPGPA CLAPQAFLQ*VLAMDKDLLEG PSWS*LGDEYNV\ESIDGQP\GA FTCYL\DADLARTTTGNKVFG\ APEGAVDGGLSNPHSSKRFLG/ YVYPLHSTQTDSPGFMDSEKQR NFNARSTSGKHIMGQ\NCLQDY IALL*WERKMKDAYKK\QFSQY IK\NSVT\PDMM\EEMYK\KAHA AIRE\NPVLWKRRPKERSLKRR GWNRSQNVPLLRKKDRVA\QR
19441	49809	A	19554	3	991	KASFLQSSRSGLLESLTPAIFP LTSRGVGSCAVMLLLVRAGAR AWLRPTGSQGLS\SLAEEAARA TENPEQVASEGLPEPVLRKVEL PVPTHR\RPVHAWIESLRGFEHE RVCLVDLHPDVFANAPRLDILH QDAMLQKNF*RISYAKTKTRAE VR\GGGGKPLAAERH\WAGPA WQHPLSALARRRCCPWPPGP\T SYYYMLPMKVRALGLKVALTV KLAQDDLHIMDSL*L\PT\GDPQ LP*PELAH\YRRWGDSVLLVDL THEEMPQSIR\EATSRLKTFNLI\ PAVGLNVHSMLKHQTLVLTLP TV\AFL\EDKLLWQDSRLQTPLY PFSL\PYSDFP\RPLPHATQGP\A ATPYHC
19442	49810	Α	19555	31	305	
19443	49811	A	19556	213	541	DGGPAAHTPGAARGELSPVGD SGGYFRPSHLRCWILNLCISVPL SSAKDLLHPSSEEE\RRNPMKN RLVQSPNSYFMDVKCPGEETV CCIGESTGPQQLENVVVLAVSY

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19444	49812	Α	19557	164	591	DINSRDRSGPRATRIDGSGEGD
						KMAAQLRRHQGRRGASSPRFA
						KD\LLHPSPEE\EKRKHKKKRL\
						VQSPNS\YFMGV\KCPG\CFKFT
						TVF*AMAQTVVLCVG/CAAPLF
						LCQPTGEKARLTEGCSFRRKHA
				1		LKATLSQDEWETILQ
19445	49813	A	19558	362	916	GGRCTMARTKADWPANRTGG
						KAPKEANWLTKSPLRKSAALY
						LEGSKKPHRYR\PGTVAPP*NLD
		1				GYQKST\ELLIRKLPFQ/RRLVRE
		ł				IAQGLLKQDSRFQ\SAAIGCFAG
		ŀ				RQVRAYLVWPFLKDTNLCA\IH
			:			AK\RVTIYAKKTFQLSTPAYRG
į		ŀ				EPCFKDPLMDGETFHSQKKKK
						KISLLPVIGSSER
19446	49814	A	19559	849	1261	DGHIAGGPCS*GWPRDPAPKSQ
						PTMSSGSEPEKLPPAPRPQHYCS
ł						P\AEVSGPRFAVGIKALNQQAFT
				ļ		ERRLHTSLCPVPAALTVPSPESL
						PQPQIFPGLAPATHTLYSKAWA
						GATVSRPGAILGAGLYWGLVR
				İ		ETAS
19447	49815	Α	19560	3	1852	
19448	49816	Α	19561	362	1298	QGGRNVRHPSPLLAWGSDLIV
						KPHWLHWLLDLNHARLSALFA
						AAGARLPSHIPAPSAGSRRLLLL
						LLGGCRACLLFFPEMANRGPSY
						GLSREVQE\KIEQKYDADLENK
						LVD\WIILQCAEDIEHPPPGRAH
						FQKWLM\DGRVLC\KL\INRLYP
						PGTRSPYPRFSESKDGFLGRWE
						QISQF\LKA\AETYGV\RTTDIFS
				}		RRWFLWEGKDMAACAEGP*L
						ALGSVAVTKDDGCYRGEPSWF
						HRKAQQNRRGFSEEQLRQGQN
						VIGLQMGSNKGASPGGP*TGYR
						ECPRQIHVRTRHPAPWLERTNV
						STPWSFTKKK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19449	49817	Α	19562	3	2518	AAGEWPQRPRAAVAAAGAAA
						RORLPGSROVAAGAPOPAGGG
						DPSQR\PPGAEILIKVQVYVSGE
						LVPLARASVDVFGNRTLLAAG
						TTDSEGVATLPLSYRLGTWVLV
						TAARPGFLTNSVPWRVDKLPL
						YASVSLYLLPERPATLILYEDLV
						HILLGSPGARSQPLVQFQRRAA
						RLPVSSTYSQLWASLTPASTQQ
					·	EMRAFPCLPGH*G\SSSGNGSW
						LELMPLTAVSVHLLTGNGTEVP
						LSGPIHLSLPVPSETRALTVGTSI
						PAWRFDPKSGLWVRNGTGVIR
]					KEGRQLYWTFVSPQLGYWVA
						AMASPTAGLVTITSGIQDIGTYH
	•					TIFLLTILAALALLVLILLCLLIY
						YCRRCLKPRQQHRKLQLSGPS
						DGNKRDQATSMSQLHLICGGP
]						LEPAPSGDPEAPPPGPLHSAFSS
						SRDLASSRDDFFRTKPRSASRP\
						AAEPSGARGGESAGLKGARSA
						EGPGGLEPGLEEHRRGPSGAAA
						FLHEPPSPPPPFDHYLGHKGAA
						EGKTPDFLLSQSVDQLARPPSL
						GQAGQLIFCGSIDHLKDNVYRN
						VMPTLVIPAHYVRLGGEAGAA
						GVGDEPAPPEGTAPGPARAFPQ
						PDPQRPQMPGHSGPGGEGGG
						GGEGWGAGRAAPVSGSVTIPV
						LFNESTMAQLNGELQALTEKK
						LLELGVKPHPRAWFVSLDGRS
		<u> </u>				NSQVRHSYIDLQAGGGARSTD
19450	49818	Α	19563	206	1126	LESGSHIYAFSHESLLLHVASAS
						V*TSFSDFATSVCHSPLHCQLH
						HALIPHGKGECSS\VSGIVATGF
						GATGFLGRCVVNHLGRVGSQIP
						QAIAQVSKEAEVGKFIHVAHLN
						VNIKSSSRYLRKKAVGEKAVR
						DTFPEAIIVKLLDIFGREDRFLN
	:					YFANMCWFGAIPLVSLGWKAV
						KQPVYVVDVSKGIVNAVKDPD
						ANGKTFAFVGPNRYLLFDLVK
						YIFAVAHSKGWVARDFEISPFE
						SWTTRDKVERIHITDMTLSHLP
						SLEDLGIQATPLELKAIEVLRLH
						HTYCWLSAEIEDVKLAKTVNI
Ь	L		·		1	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19451	49819	A	19564	196	553	KLSSEFFLTHSINTAIRNGNLFV
						NRVTEVTRSGGNELGSFCLLAI
						KFTLLSFLGTGTRRLLPALRPRR
	1	ĺ				LQQRGSHGLAGRLFTLWILHA
	-					GCHARGELLEPDTC*LSLCRAR
	ļ					KPRLSTLK
19452	49820	Α	19565	1	505	ATEDPGVAMGRRPARCYRYCK
						NKPYPKSRFCRGVPDAKIRIFDL
						GRKKAKVDEFPLCGHMVSDEY
						EQLSSEALEAARICANKYMVKS
	'					CGKDGFHIRVRLHPFHVIRINK
						MLSCAGAD/RIHISKKWGFTKF
						NADEFEDMVAEKRLIPDGCGV
				İ		KYIPSRGPLDKWRALHS
19453	49821	Α	19567	6	904	AQLSSLAVPLLVLTPFRSVLVFR
						SDSLFFVAATEDPGVAMGRRP
		İ				A\RCYRYCKTKP\YPKSRF\CRG
ĺ						CPLMPRIRIFD\LGRKKGKKWN
				1		EFPLLCP/HMVSDEYEQLSSESP
						GRLPRICPNK\YMVKSCGKDGF/
ŀ						HISRVRLHP\FHVIRIHKMLSCA
						GADKAPKQACRGAFWKSPKGN
						LLARGSQLAQVIMSIR\TK\LQN
						KEHVI*GPAARPSFKFPGRQK\I
İ				1		HI\SKKWGF\TKFQC*WNLKDM
						VAEKRVHPHMACGGQSTSPSR
						GPL\DK\WRALPLHEGLPNVAA
						HPLNTPPIKFLLPWSKKKKKK
19454	49822	C	19568	346	471	
19455	49823	Α	19569	1495	2780	EGLHLFCDLLDGLPGVQVAGG
						FDSFSSILASFTY WHDFPNSTML
						LSTFRDINLGMNIIKKSAVCFSR
				}		LGSETWEMPANLRPSEGGART
				1		RTELSRPQTPSRAHLAWRGRGG
						LGAVRPRRLSRRRSLS*KP\RSL*
						AEAPAGAPVRRCRPAAPP/PPRP
						SRCGPSTPPPPPSPGA\PRRRPP
				1		CSEHNKLSPPPPSSLCLRPAPSPP
				1		RPAPRRSPSRRRRRRRHPRPAD
						RREPLASATRVRPLPPPPPPR*PP
						SSSCPRRGGRRFLAPWLARPAG
						APRRAPCSRAARSRRAGASGAR
				1	:	VGDPAGAGRGWRRRAAGRP*
						APERRPPPPLPAVCEVSGGPLRA
						SFADSVPGVRRRRPPGLGTPLSS
				1		SGV*PRNERGVETTTSWGGGSE
						GAPGSGEGGNLFPAPQVIVARS
						GEGTVVPRFPVCQLRDSAVLLC
						LFD
19456	49824	A	19570	178	256	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19457	49825	Α	19571	622	1726	QGSGRTMMRMLVPASVGLLET
						GPPGDPSLSHPLSGDVTVLPED
						Q\STVQKLASDHKDIHSSVSRV
						GKAIDRNFDSEICGVVSDAVW
1						DAREQQQQILQMAIV\EHLYQQ
						GMLSVAEELCQESTLNVDLDF
<u> </u>						KQPFLELNRILEALHEQDLGPA
						LEWAVSHRQRLLELNSSLEFKL
						HRLHFIRLLAGGPAKQLEALSY
						A\RHFQPFARLHQREIQVMMGS
						LV\YLRLGLEKSPYCHLLDSSH
						WAEICETFTRDACSLLGLSVESP
						LSVSFASGCVALPVLMNIKAVI
1						EQRQCTGVWNHKDELPIEIEL\S
						*SAWYHSVFACPILRQQTSDSN
						PPIKLICGHVISRDALNKLINGG
19458	49826	В	19573	271	392	KLKCRYCPMEQNPADGKTH
19459	49827	В	19574	58	296	
19460	49828	A	19575	215	861	QLPAESVRTRGPAKAQARAGG
17400	77020	 	17575			PPGPWHCSYKWRAWPSSLSGT
						GWPDMEGETTAAAPASRENSA
						DCS\AGELRGPG\KELW\SPIQ\Q
						VTATSS*MGRDFVLAT*EKVSH
j						VDS*/EQPRSLQR\DPRP\AGPAQ
						AKQGTPRAQASRESL\SRPTAA
						V\QLPSGPQHPV\TSGS*KGLAG
						KTSMGRKGLPWARGWAPWSL
•				ĺ		EAQNP\RPTRLCDLFITGEDFDD
						DV
19461	49829	A	19576	65	782	SSSSSAIQRKPRWVLYKYIQEL
15401	77027		17570	05	7.02	WRKKQSDVMRFLLRVRCWQ\Y
	İ	İ				RQL\SALHRAPRPTR\PDK\ARRL
						GYKAKQGYVIYRIRVRRGG\RK
				ŀ		RPSFLKGAT\YGQSPVHSWVFN
ŀ				ŀ		QLKSFASE/VLSVPFARGRRSLG
	1					RH\CGGLLEEFLNFFNWVWVER
				ŀ		FPHYQIFLKVIPHLIPFP*KLFRR
		1		}		N\P\DTQLIHQNQVHKHR\EMRG
1		1				\LTSARPEKSRGLGKGHKFHHTI
l		1				GG/SLRRATWRRRNTLQ\LHRY
						R R R R R R R R R R R R R R R R R R R
10462	49830	A	19577	1	179	QEDYLRELLTTMGDR\FTDEEV
19462	49000	A	19311	1	1/3	DELYREAPIDQKGG/MFNYIEFT
		1				RILKHGAKDKDGLKEL
L	<u> </u>		L	<u> </u>	<u> </u>	MILMIUMNDALKEL

SEQ ID		Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19463	49831	A	19578	144	722	KKGLNHHHVEQNNTRPRPRSG
						PQRATSN\VFAMFD\QSQIQEFK
	1	1				EAFNMIDPDQDGFHRPGKICND
						MLASLG\KNPT*WSILDAMMNE
		ł				\APGPPSNFTPWFPHHGFGEEV*
						MATRS*KMVIRKCPLTCFE*KK
		ŀ				PLGTI\GEDYL\RELLTTHGGIGF
						TDEEVDELYREAPI**KRGISITS
						EFTRILK\HGAKDKDD
19464	49832	Α	19579	3	429	VACLREHKQRCERATRSLLREL
						LQVRARVQLQGSELRQLQQEA
		İ				RPAAQAPEKEAPEFSGLQNQM
						QALDKRLVEVREALTRLRRRQ
E						VQQEAERRGAEQE/DRPQTGQA
						DRLAAAGGAGPGGGLRRPAEE
ļ		1				PRGQQPEGGPGGGQN
19465	49833	Α	19580	1	2454	RFPAGVAPRRAMANVSKKVSW
						SGRDRDDEEAAPLLRRTARPGG
						GTPLLNGAGPGAARQSPRSALF
						RVGH\MSSVKLDDELLEP\DMD
						PPHPFPKEIPHNEKLLSLKYESL
						DYDNSENQLFLEEERRINHTAF
						RTVEIKRWVICALIGILTGLVAC
						FIDIVVENLAGLKYRVIKGNIDK
						FTEKGGLSFSLLLWATLNAAFV
						LVGSVIVAFIEPVAAGSGIPQIK
						CFLNGVKIPHVVRLKTLVIKVS
		ļ				GVILSVVGGLAVGKEGPMIHSG
						SVIAAGISQGRSSSLKRDFKIFE
						YLRRDTEKRDFVSAGAAAGVS
						AAFGAPVGGVLFSLEEGASFW
						NQFLTWRIFFASMISTFTLNFVL
						SIYHGNMWDLSSPGLINFGRFD
						SEKMAYTIHEIPVFIAMGVVGG
						VLGAVFNALNYWLTMFRIRYI
						HRPCLQVIEAVLVAAVTATVAF
						VLIYSSRDCQPLQGGSMSYPLQ
						LFCADGEYNSMAAAFFNTPEKS
						VVSLFHDPPGSYNPLTLGLFTL
						VYFFLACWTYGLTVSAGVFIPS
						LLIGAAWGRLFGISLSYLTGAAI
						WADPGKYALMGAAAQLGGIV
		1				RMTLSLTVIMMEATSNVTYGFP
						IMLVLMTAKIVGDVFIEGLYDM
		1				HIQLQSVPFLHWEAPVTSHSLT
						AREVMSTPVTCLRRREKVGVIV
						DVLSDTASNHNGFPVVEHADD
						TQPARLQGLILRSQLIVLLKHK
19466	49834	В	19581	508	4413	

SEQ ID	SEQ ID NO:		_		1	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
		<u> </u>				
19467	49835	Α	19582	8	331	EVTACWQSSQPSLTLSASSAWA
	1					STLAALEEPFSPPLHSESPFLGW
						PRPELGPSAAPCAPVPPEPPQRV
						PPPAP*RPVPSTTQGLRSVGAW
		<u></u>				CRTGRQLHLHPGVGSTG
19468	49836	Α	19583	831	1121	
19469	49837	С	19584	1	1524	
19470	49838	Α	19585	270	578	RFSKSPPDSGAQLAAPSGSRTG
						AAGGAACQSRAVRPHSSALGW
						SMGLGAMEQGAVLVGEAWAT
ŀ						QQPTVGGRLRHGGLQVPSPAPR
10471	10020	<u> </u>	10506		200	GGS*GPAKSHSSCWPRC
19471	49839	A	19586	2	298	
19472	49840	В	19587	113	515	MI DONI DONI A MUDOD A CONTI
19473	49841	Α	19588	1	569	MLPDRLRSPVAHHRCRASGNY
						HSGRHSPAPAWRVTLTGWLDQ
						RSSRIHRSLAFKDSYSKGKEERT
	,					HIKGMCSELLPSG/EARGLAGFR
						SEAADLRGECYSS*RQCGPKEL
						RSPAGFTQWIPHWGYRWSCLS
						VPRPAPALLSPWVVDGTGRCG
						AGGGAHRGGSGRSGAHGGGG
10474	10040	<u> </u>	10500	1	1200	RLRHGGLQVPSPAPREGS
19474	49842	C A	19589 19590	511	1200 846	ARHRVLIGVFTIPELDIKVLHVP
19473	49043	^	19390	311	1040	TRLRSPASFTQWIPHWGCRWSC
						LPVPRRVPALLSPWVVDGTGG
						CGAGGGAHRGGWGCTGAHGG
		ŀ				GGRLRHGGLQVASPAPREGS*G
						PARN
19476	49844	A	19591	9	481	QQPLREKPASGDHPAKNANRR
17470		` `	17371	ľ	101	WPSPTRYPYVES*NWMPSLHLI
	1					QKSIQDGLKT*TLDLKP*KL*KK
						T*ALPFRT*A*ARTSCLKHQKQ
						WQQKTKLTNGI*LN*RASA/PA
						KETKI/SSEQATYKMGENFHNL
						LF*QRANIQNLQ*TQRDLQEKN
						KQPHQK
19477	49845	A	19592	1	3339	MSVRKDVEKLEPSDIVCGNVQ
						CYSCMETNLTVSQKVKHEVTV
						GPREGATKPNRMKGKEGRSGS
						LLGEGDFFKDESVMSSQGSSKD
						GEKRRGKAQRWKWPMQGICR
1						QLGVAKSMEGYQSRRDQGGR
						GVSDKWPQVCAKKPEFYPTAQ
1			}			VWANFSVTSCQSVTITQLCHGL
			1			RRLEISPARSNAMHLNPDPPGQ
						KQNLSPKVNDIITDIESSSGSGA
						GKFQVISKSDISEVLLQQMDAG
						HSSKDDPNEYGGWKSPRPRC
19478	49846	В	19593	1	631	
194/8	47840	ĪΒ	19393	11	031	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19479	49847	Α	19594	71	208	PCPKHCSKFPMCPELVGSWSH*
	;					LQE*SHGPSRFRLC*KQPQDLQ
19480	49848	Α	19595	138	650	AWAQLPSALGSTRVSPPPSARS
						QPPWRRRHKVCWRLNRCKE/L
						DEDAVASADFSSMLSEEEKEEL
						KAELVQLEDEITTLRQVLSAER
		l				KAS/ELR*NKNSA*T*/CNELKPE
						LQQKLA*HAD*HCLQEDT*YAE
						SRRAKGNCSFQQRWNGHQQEV
						RRHELLHSPFHKYACYETKVG
19481	49849	Α	19596	1	1222	MRADGTAGKDPFATDKQLPEL
						LIQHCWDGLLAVCGGSTVGVR
						LPFILHGSWVRPVAASFPPLPDN
		i]			LHDSAEPAIILLEPTQMRRNQK
						TNAGTMTKQISSTLPKNHHTSS
		ŀ				PAMDPNKEEIHDLLEKEFSRLVI
						ELIREGPEKGKAQCEEIQKKDT
						KVKGEIFKEVDSLKKKQLNIQE
						NLETLLEVLNALESLSNRIEQVE
			:			ERNSEREDKVFELTQSNKDKA
						QRIRKYEPSLQEVWDYVKLPKL
						RIISFPNEEENSKSLENIFRGRIK
						EHFHSIARDLDIQIQEEQTTPGK
	:					LIVKRSLPRHIVIMRAPTPMDHP
		ļ				RAEECGRRARDWQAAPPAAPV
						RDPLGEASWAPDSEGRK\WNTS
						KHQRNKLRTRHL*EL*HSSRGS
						VASFLKSVRPRTHQFQTQSGYP
						KLRWRKES
19482	49850	À	19597	381	598	
19483	49851	Α	19598	1	1906	
19484	49852	Α	19599	493	589	
19485	49853	Α	19600	60	261	
19486	49854	Α	19601	1423	1831	PRRSANVNLPPPQLATLSPGWP
	1	l				EERKSVGSPTSPPPPRPLRRLSA
						APG\SGGAWPEPGAWSLD*TAS
						PAGAPVQRPAGNG/PSALDASH
						ARALYLQRIPRVTWTPASQPPA
						PLGPFLPLIPPAASHRQPSSSFFR
				1		RHR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19487	49855	A	19602	1	1535	MEEKTNVQLPPGQTEQHVEIHI
						MNFCSKNHHRITPEKPKELTDP
						FKEAACCCKLYEIDKKLYRMA
						EWIKIHKPSICCLQETHLTHKDS
						HKLKVSITFKDLAVRFSEEEWR
						LLEEGQREFYRDVMRENYETL
						VSVEPGRAVGGGSHADEGQEP
						AGCG/VSPGPGAAGEGDPRVLV
						WRSQGRYGQPRER\GRGASLD
				•		GERASPEAA/DGKRALPSPRPA
						QLPSRRPYQPAPPG\PTPTDSSCS
						SGPTGDGVQGSPLPIRISPGNSP
						L/PRPHQLSEGNPCAWAPAPRDI
						PKLLATSP*PGHVQANQSRPGA
						WEPALGRSDQRACSASGSAELC
						ERWPQQAP/APPEEPPPASPHPA
						APTG\PGFWESCGEPGAA\PGKG
				+		SAPKPSPLHCLESALRGILP\EGP
						CASPAWEAPAPAPAPAPARASA
						A/AEGEDPRPEPELWKPLPQER
						DRLPSCKPPVPLSPCPGGTPAGS
						SGGSPGE\APGEQSPGTAAASV
						Q/VSPAHWPCFS/SPVRYSSGSLP
						GFSAGEKAQG
19488	49856	С	19603	4	150	
19489	49857	Α	19604	148	448	RTRDWSLFLLMFGCVRSFFLLV
						GLWSCWLRSEAADLRGECYSS
						*GGASGVVRSSWWVRGLTSFR
						SETGFTQWIPHWGCRWSCLPVP
						RCAPALLSPWVVNG
19490	49858	Α	19605	136	439	LVHFTEC*LVRLQTLARHSTDW
						CIYKSLARQKSSPSPHSTQEVQL
						ASPLTGRQEDCSEPRSAPLHSSL
						GPTERDSICVAASSVRWTRPGA
						RFS
19491	49859	Α	19606	2	399	SKLQKLKRSLSFKTKSLRSKSA
						DNFFQRTNSEDMKLQAHMVAE
						ISPSSSPLPAPGSLTSTPARLVCI
						QVARLMPFRNTSSRSPLFC/RCL
1						QPHDSGFLFLRPARPRAHREER
						NNSRRVALRAVTLTTERLQLHS

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19492	49860	A	19607		1262	MPLSYAYKNAETLAGRHTSSW MSRGAYQRRNTRAAGRPEECT DRNWHAGRTRRIWLGQLEERC SDVFGVSFFWWVRGLAGSGAK LQTFTVSVTAHRGSKDQKSEQ KQDLLQKAKEQTIHNVKGDPS ELPLLAPAACFYSLICPRPPPHIL LIGPFYRELIGPFYRELIGRQLRP GEKSSTAA/VWPRC*AA/RLPGA G/WTAPSAGPAEPTPTWNSRWP ASATRRPGSRRRLSLHTSPQAE GAGSGLSQPRKGLPQCSGRLKR SSSAARVGVARQRRRERASEG/ WPACCHLSVLPCGFGFLEAHSS SPEQTDHEQNCPADIKVMRLA GQPTNARGAANRAQPLHSTSV SHLKAVIFKIQTRQQKNSHFPPL FLEMTTSLATGRLRPSWVDYA RPALSVVSVSRFSLLTSLLELLV ETLGECGLL
19493	49861	C	19608	173	394	
19494	49862	Α	19609	1	924	
19495	49863	A	19610	3	689	EVQEELEGPVRGAPHQLHQGP VEHLEAPHQ\HKFGKKVIYFNY LSELHEHLKYDQLVIPPEVLR\T MRSSRACTRAGRRLPPRHHRR GPRCPHSSLASVCNTSKTKIKA NSSPLC*GSQ*RT*ERKACAPRA CSGDPPACRPSARSRGSTT/KGK PVNFDDYGDIHIPAVILKTFLRE LPQPLLTFQAYEQILGITC/GGE QPACHSLPPDLTEPPRAQLRRPP LPHGLPACG
19496	49864	A	19611	1749	2411	APSCLVSEHSAPGPQRELPQPLL TFQAYEQILGITC\GSCP\AQGW GAWSSDAVPQLLARRPPLPHGL PACGEWGRGELGVKPSGLPSH AGPAWGHQVRTVCATAHPQD CISPEGAVEEEIVGG*GC\TEGQ SQRVLQIWP\SQGVSSLSALVPL N\MF\TELLIEYYEKIFSTP\EAPG EHGLAPWEQGSRAAPLQEAVP RTQATGLTKPTLPPSPLMAARR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19497	49865	A	19612	883	2164	EHRRPMRAQASHEGHRHPMRA
17477	17003	1	,,,,,,			QMSHEGYRRPMRVQASHENTG
						VP*GVQASHEGTGIP*EHRCPM
						RTQVSHEGTGVP*ECRRPMRTQ
						\GPMRGTGVP*EHRCPMRGTGV
						P*EAQASHEGHRRPMRGTGIP*
į						GAQASHEGHRCPMRGTGVP*G
						HRRP/DENTGVP*GHRRPMRAQ
						ASH/REHRCP/DEGHRRSP*GHR
						RPMRAQASHENTGVP*GAQAS
		:				HENTGVP*GHRRPMRGTGVP*E
						l .
						\QGVP*GHRRPMRAQVSHEGHR
ŀ						RPMRTQASHEGHRRPMRTQAS HEGTGVP*EHRRPMRGTGVP*E
						l .
	:]				FRRPMRGTGIP*GHRRPMRAQA SHENTGVP*GNR/PYHEGTGVP*
		1				
		l				ECRRPMRTQASHEGTGVP*EHR
						CPMRAQASHEGHRHPMSTQAS
						HEGHRRPMRTQASHEGHRHPM
						RTQASHEGT\ASHEGHRRPMRT
		1				QVSHEGTGVP*EFRRPM/PGHG
ŀ						HPMRTQASHEGHRRPMRVQAS HENTGVP*GHRHPMRGTGIP*G
						HRCPMRGTGVP*GYRHPMRTQ
		j				ASHEGYR/PSHEGTGIP*EHRCP
		Ì				MRTQVSHEGTGVP*ECRRPMR
İ						TQVSHEGTGVP*GHRRPMRTQ
						VSHERHRRPMRTQASHEGTGV
						P*EHRCPM/MGHRRPMRTQASH EGTGVP*EHRRPMRGTGVP*EH
						RRPMRAQESHENTGVP*GAQA
						SHENTGVP*GHRRPMRAQASH
10.100	10055	<u> </u>	10610			
19498	49866	Α	19613		552	VPQDSLGQALAGQATPEIPLGL
						QLHTVLVQEIQELIEAQTRAPGP
						CAEVRALPAPAAEPEPAWEEAP
						PERALELEGAPAKDQTNEELPEI
j						TA/PYC/EPLALTLELKAWLERE
						VGGRG\DQHSPSQQLPCCP*SW
						ARWQTCRQRAGHLAWPPVPRC
						REASLIH*NHSPAAAGPFILLILV
10.100	1006	ļ. —	10514	1.12	001	YLCAGGA
19499	49867	Α	19614	443	991	VDSRPMLNTRLMRTSGRRWTP
						RPQVDTRP*VNNKTQ*AIRPQ/C
						GYSPQVPQDSLGQALAGQATP
						EIPLGLQLHTVLVQEIQE/RH*G
	:					TDTCSWSMCRSKSTASTSRWS
			}			Q/RPAWEEAPPER/GTGAGGSSS
						QGPDQRGAA*NHGGAGEH*KK
						AGRKSSE\RYACSQRQPPPPGM
		_			1.50	DAGPQLGIQAVCP
19500	49868	Α	19615	16	450	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hođ	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
İ						
19501	49869	Α	19616	2	477	PRVRSGHSMDMRVPAQLLGLL
ŀ				İ		LLWLRGANCDIQVTQSPDSLSA
ļ						SVGDKVTITCRASQGISNGLSW
						YQQKPGQAPTLLIYAASSLQSG
1						VPSRFSGSGSGTDFTLTISSLQPE
		1				DVATYYCLQDYTTPFTVLQAIT
1				j		KTPQR\RDV*VCTVLATPPGVIN
		l				WL
19502	49870	Α	19617	2	675	LPLSSSALPTESGSHLPFSGIMA
						RPSDPLVKPQRSSKREPKKVHP
]						GTQSDR\YVKIKR\NWAENPRG\
İ						IDN\RVR\RRF\KGQDLDAPTLG
ŀ		1				YG\SNKKTKAHACPSGFREVPG
ŀ		1				VHNVQGSWEVACLMLQQNLY
						CCRGSVHNVFLPRNRQSPSWG
						KELAQLGLSGVHPTPLARAAQL
						KEN*VRQLMCNKSYCAEIAHN
						VSSKNRKAIVERAAQLAIRVTN
		ŀ			:	PNAR\RSEENE
19503	49871	Α	19618	2	527	VWAANMPSRLRKT\RKLT\GHG
						AATGHGPHRPSTRKAPPGGPR*
						MPGGPAIHPPGSNFRQNTHPQA
						YFGK\VG**KHYPT*KRN\QSFL
					1	PPTVNL\DKLWTLVQ*NRHRVK
]		İ	CLPKNKTGAAPHHLMLVRSGY
					•	LQSSGERGKLPKASLFFVEGPK
						FFSRRSLREEGLRGVWGGPVSL
19504	49872	Α	19619	45	233	
19505	49873	С	19620	95	154	
19506	49874	Α	19621	2	241	
19507	49875	Α	19622	63	320	RQWGNRLVTA/WSFFSWLSDC
		1				EGPGYGTIAEVGQLWGIMAKD
		l				LLRHSLASEELALNQNLTCHDQ
						FAQSTRSSARACGSIFCSTTQGS
19508	49876	A	19623	2309	2560	RNREARFYMTTPNFTETKR/WF
						HERVCALEREREREVTKISRLE
						RDNHRLVGDISQLKKELDQYL
		L_				QAISDLEDCNGKRLLQILEA

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19509	49877	A	19624	4994	7008	QWQKPKLTNDNLKFIWNQKR DRIAKTILSQKNKAGGITLPDFK LYYKATVTKTAWYWYQNRHI DQWNRTEPSETIPHIYNHLNFD KPDKNKKWGKDSLFNKWCWE NWLAICRKLKLDPFLTPYTKIN SRWIKHLNVRPKTIKILEENLGN TIQDIGMGKDFMTKTPKAIATK P\N*QMGSNILNGQKLEAFPLKT GTRQGCPLSPLLFNIVLEVLAR AIRQEKEIKGIQLGKEEVKLSLF ADDMIVYLENPIVSAQNLLKLIS NFSKVSGYKINVQKSQAFLYTN NRQTESQIMSELPFTIASKRIKY LGIQLTRDVKDLFKENYKPLLK EIKEDTNKWKNMPCSWVGRINI VKMAILPKVIYRFNAIPIKLPMA FFTELEKTTLKFIWNQKRARIA KSILSQKNKAGGITLPNFELYY KATVTKTAWYWYQNRYIDQW NRTEPSEITPHIYNYLIFDKPGK NKQWGKDSLFNKWCWENWLA ICRKLQLDPFLTPYTKINSRWIK DLNVRPKTIKTLEEDLGITIQDI GMGKDFMSKTPKAMATKDKID KWDLIKLKSFCTAKETTIRVNR QPTKWEKIFATYSFDKGLISRIY NELQQIYKKKTNNPIKKWAKD MNRHFSKEDIYAAKKHMKKYS SSLAIREMQIKTTMRYHLTPVR
						MATIKNSGNNRQ

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19510	49878	A	19625		764	GSSEKRKFKTGNQKLSMEVEK NQQMRSLSLYQQRVEGSKCAV RRKGGSPKPKAFLAAAEKFLN VPVMRSASHEQLVLTGDPNKT LPGSKSRLSGTDSPRACSTFPIW ATKIVTEMQVQLFAACRVQLIK HPRPSAARTAMEVLESGEQGV LQWDRKLSELSEPGDGEALMY HTACQGHEEQVWEEEALQLGL RSGILLNHWKRQLVGGAGILRS AQLHSPYHHWQMISLVSPSPAV MAVMENDDICMVLCVKVMDE PNHILDAISEHQIPSLEDQWFSE CALWTSKAYENLVERHILGPHC RPTELETLGISEDDIDHNPYLDF PGVAGLGSRRNVKERGWLSEM KKEGDIFMKIQG*AICPTSGPAE GLPPPTPVPHVPDRPCGT*GP/PP SPGSDSS\PACGPTAARPAPRSP APPWRCGPRWAEDVLLTGLCK QQIAEPAFRLQFWWPI

2457

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19511	49879	I _A	19626	<u> </u> 1	1430	L MNGTAEGQHPWVHSTFNGEPQ
19311	49879	A	119020		1430	SLSSSVRVKNQGPRDQAELGFQ
		1			·	GNGSTVPAAQARTIARARGSGS
						GGGDALQGGGAGAGPGATKR
						TARLLARGLRRAALRPDGRGV
						LGAGAGSSQRRGGGRRKMETH
						ISCLFPELLAMIFGYLDVRDKG
						RAAQVCTAWRDAAYHKSVWR
						GVEAKLHLRRANPSLFPSLQAR
						GIRRVQILSLRRSLSYVIQGMAN
		ŀ				IESLNLSGCYNLTDNGLGHAFV
						QEIGSLRALNLSLCKQITDSSLG
						RIAQYLKGLEVLELGGCSNITN
						TGLLLIAWGLQRLKSLNLRSCR
						HLSDVGIGHLAGMTRSAAEGLP
1						GPGAAHATGLPEAHRSFSKAHL
						PRADGPEAPQPQLLWGNLGRW
						PPAPVAHGQPAHAQPCAPCDNI
						SDTGIMHLAMGSLRLSGLDVSF
		İ				CDKVGDQSLAYIAQGLDGLKS
						LSLCS/RRGLRRAALRPDGRGV
						LGAGAGSSQRRGGGRRKMETH
						ISCLFPELLAMIFGYLDVRDKG
						RAAQVCTAWRDAAYHKSVWR
						GVEAKLHLRRANPSLFPSLQAR
•		1				GIRRVQILSLRRSLSYVIQGMAN
						IESLNLSGCYNLTDNGLGHAF/R
						AGDRLPARSQPEPLQADH*QQP
		İ				GPHSPVPQGP/ARCWSWEVAAT
						SPTLAFCSSPGVCSASRALTSAA
						AATFRMW\DRAPGRHDAQRGG
						G\LPGPGAAHATGLPEAHRSFS
19512	49880	Α	19627	293	1520	TTMDGSHSAALKLQQLPPTSSS
						SAVSEASFSYKENLIGALLAIFG
						HLVVSIALNLQKYCHIRLAGSK
		l				DPRAYFKTKTWWLGLFLMLLG
						ELGVFASYAFAPLSLIVPLSAVS
						VIASAIIGIIFIKEKWKPKDFLRR
						YVLSFVGCGLAVRG\TYLLVTF
		l				APNSHEKMTGENVTRHLVSWP
						FLLYMLVEIILFCLLLYFYKEKN
	1					ANNIVVILLEVALLGSMTVVTV
						KAVAGMLVLSIQGNLQLDYPIF
						YVMFVCMVATAVYQAAFLSQ
						ASQMYDSSLIASVGYILSTTIAIT
	1					AGAIFYLDFIGEDVLHICMFAL
						GCLIAFLGVFLITRNRKKPIPFEP
						YISMDAMPGMQNMHDKGMTV
						QPELKASFSYGALENNDNISEIY
						APATLPVMQEEHGSRSASGVPY
L	<u> </u>	1	<u> </u>	<u>L</u>	<u> </u>	RVLEHTKKE

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19513	49881	A	19628	751	2034	VSSSPTSLGCMRQKEYRIPKKL TNMPFLGSQPSLLALRSCHPTG RTCSA/SQVALFCLAAHRVSQIG ALLAIFGHLVVSIALNLQVSFHI RLAGSKDPRAYFKTKTWWLGL FLMLLGELGVFASYAFAPLSLI VPLSAVSVIGKTIIGIIFIKEKWK PKDFLSKFVLSFVGCGLAVVGT YLLVTFAPNSHEKMTGENVTR HLVSWIILFCLLLYFYKEKNAN NIVVILLLVALLGKLTVVTVKA VAGMLVLSIQGNLQLDYPIFYV MFVCMVATAVYQAA*VSQAS QMYDSSLIASVGYILSTTIAITA GKGFYLDFIGEDVLHICMFALG *VLAFLGVFLITRNRKKPIPFEP YISMDAMPGKVNMHDKGMTV QPELKASFSYGALENNDNISEIY APATLPVMQEEHGSRSASGVPY
19514	49882	A	19629	19	933	QSARPSLKRARSQRGRPLPSRA LVFLFLFFFFFFNFLPIHTQPSAH KDMTTNAGPLHPYWPQHL\RL DNFVPN\DRPTWAYTGLGLFSV TGGLRSWTTWAVVQGRAPGL\ PIGDFGRATVPCCWVLQWCGFI HLVIEGLVPFSNYERPCFGDQSL LYSQLW\KEYGQGETARYILGV TNFTV\CM\ETITSFA\WGEPFSL WVVIV\FLLQHPL\RFIL\QLVVS VG\QIYG\DVLYFL*RATPTDFQ HGKPGPTPLLLPGFSFVFHEMP LWLVPAPGSPLCLDA/V*KHPQ LMPQSNAWNAKGPQKPKSKK
19515	49883	С	19630	113	257	
19516	49884	A	19631	143	330	HHGPVPFPRGLIS*HSLHCPCPT PGGKNVFFFCEYFLKLAGIETT NCSADKRGKKVCCLIPR
19517	49885	Α	19632	3	432	
19518	49886	С	19633	112	192	
19519	49887	A	19634		4410	MARPLALPLTGMSAPGSSLTTIP HTPSHPRSPSTPDTVSHPRSPSTP DTVSHPRSRYTPDTVPPTLPVY TRHRVPPTLPVYTRHRPTHAPG IHQTMCPTHAPGLHQTPCPTHA AGLHQTPSHPRSPSTPDTVSHPR SRSTPDTVPPTLPVYTRHRVPPT LTVYTRHRVPPTLPVYTRHCPT HAPGLHQTMCPTPRSRSLPDTV PPTLPVYTRHRVPPTLPVYTRD RVPPTLPVYTRHRVPTHAPRLHQ TPSHPRSR

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19520	49888	Α	19635	1	444	
19521	49889	A	19636	3	1017	SSRPVRPRPAARLSAMSSTQFN
	1					KGPSYGL\SAQ\VKNRLL\SKYD
						PQKE\AELRTW\IEGLTGLSIGPD
						FQKG\LKDG\TIL\CTLMNKLQP
						G\SVPKINRSMQN\WHQLENLS
						NFIK\AMVSYGMNP\VDLFEAN
						DLF\ESGNMT\QVQVSLLALGG KRPKTKGAAEGGLDIGVKYSE
		İ				KQERNFDDATMKAGQCVIG\L QMGT\NKCASQSGMTAYGTRR
						HLYDPKNHILPPMDHSTISLQM
						GT\NKCASQVG\MTA\PGTRRHI
						YEYQAGNPTSCDNFSM\SLQ\M
						GYTQGAQTQSGQVF\GPGRPDI
						*TPSTCPQGTI\ADGAPSGTG\D\
						CPDPGEVPEYPPYYQ\EEAGY
19522	49890	A	19637	1	479	MQGLRVSHSTFYTLIPEVTENP
	1,50,50					HRLDFHWFLSAQLERVASGTIL
						KKKESPPRPKVLGSARWDLPV/
						PPGWCNSTALQ*SPRSDQV*SE
						RGMEAFFLPMPSESVASVCCRF
						SPWTSQKPQLSTSRDRKLQGKR
						*WFSTGVWAS*DHS*ERRCSQC
						PKTTCW
19523	49891 -	Α	19638	2	501	FVPEERSSETALQRCSLPPGRRA
						APDPGLQGLRGSLGGSGSELES
						RKAKLRAAEGLRRKTTEINQRD
						TADRCTRGHH*SPGESPPKPSAF
						SL*LSVPRSGKLRFLRSPRREPA
						AN*SHRLRRHREKKCFHPPFTL
						NLIASGTSLEGGGVAPTWSTGR
						SQRALPSTL
19524	49892	Α	19639	$\lfloor 1 \rfloor$	1122	RFLILINHAVFEYQDIFVASFPSI
						FFNIHCRGEGEREKGFAGKDRT
						PPPSTLLCGSGHLPPLASRRPPR
						PRCACCLPGGPLSSSALEAG*Y
						WRFPKLSG*KRSHTPWVPQPQQ
						PLPNPSPGKWVWASSEALQ/GT
						RA*RGIAELTHLLRLPARRGTQ RPA\PGHRAGASPQFRREASSRG
						PGLRPRPGRTRELGFPKRRVWA
						TAASWPRGTEPCAAA*RRRRRR
						RRARGRELSPRSTWSSDTESTF
						ALSCDPS*PRHQSVPKGRVGSC
						KGKLANKGFFCPTWQRPIRVAS
						VDMELLHFVPKRALPGFTIAISF
						LSSAPPLCPWYSGSNPWLRNPL
						KPRLAALFQRSAAPRLHFSGAA
						,
	l			<u> </u>		SLQVGERRQIRDFRVSAGA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19525	49893	Α	19640	14	618	GNENVKSGRKTSLSPFVLSDFT
						ASSALLASDSSRWMSLQSASEC
						\PGRAAGSGPGYG*PVSEVTGG
		İ				PLSAHTHSSL*TSQRGCPPG\PE
						HHGAGPFP*GPGLKDVSGRGG
						NGCWGRRKPAHMGGGLGPAT
						PLQFPPRPSPPPPPLGPPQ\PPPPP
						SPPLGPASSPP/CLPHRPPPPPPPP
						PSRCVTAPCVTAPGPAPTGAAA
						RRH
19526	49894	В	19641	42	121	
19527	49895	A	19642	1	153	
19528	49896	A	19643	104	355	DDDTMAIL AICIAN LLTVI OVCD
19529	49897	A	19644	459	988	DPRTMNLAISIA\LLLTVLQVSR
						GQKVTS\LTA\CLVDQSLRLDCR HENTSS\SPSKYEFSLTRETK\KH
						\VLFGTVGVPEHT\YRSRTNF\TS
•				-		KYNMEGPLTYSAFH*ARTRGTL
						QVLQFHQFGALSPPIFLSRNV\T
					·	VVQRTNWVKC\EGI\SLLAQNTS
						WL\LRLLLS\LSLLQ\ATDFMSL
19530	49898	A	19645	1	950	WE LET CE EST CONTROL WITH THE EST
19531	49899	В	19646	55	1459	
19532	49900	A	19647	1	165	
19533	49901	Α	19648	183	700	PNGNQPPRETPQRISRGIQRLGT
						AVASSGGSRSLPRSASAEPAQG
						AAARRAAPGPAGRCRAARGAG
						AGSARRLP*APAHPAPPEQGRR/
						SRTARSRAAPQPPPPSQPQSPQL
		1				GPRPSQGAPQRRGPSSPGPRAG
						CRAQ*EALTPRLWRSSRTPSSR
		<u> </u>				GLGPHVRGGIFSPKIPT
19534	49902	В	19649	71	264	
19535	49903	Α	19650	3	810	GVSPCWPGWSRTPDFGSNPKCP
}						PIRASPGAELQALSSTVTTPYW
						GILVTAVFPH*GLRPRQCRQDH
		1				PAGRQGPGPGEVPEILGQSGCT
		1				DRTWSKAGGRTQAPGPRSRAG
						RRVSGQEIRAPGPLGCRHGG/V
						GAPWTPEAASPLTATEPSCPH/L
						QAPCGYMPLSVSPRRRYRGPA
						GDQKVKMLKFKAFCLDYWQF
						LCLQPLHGAYKRDSDLMTWIW
						GLLPEVTGAAGTTSPNVHTSGR
						FFRACVFCPVHTLVKKEPHPGQ
L	<u> </u>	<u></u>		L	l	QEIIMEPSPWSP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
1				sequence		
19536	49904	A	19651	258	1378	REILSLDRLPGATPNQALSAEA
1,7330	17704		17031	250	1370	ACPKGQTSLLCGAMGIQGLAK
						LIADVAPSAIRENDIKSY\FGRK
						VAIDASMSIYHFLIAVRQGGDV
						LQNEEGETTSHLMGMFYRTIR
		l				MK\ENGIKPI/YKSGELAKCSEL
	;					R\AEAEKQLQQAQAAGAEQVV
						EKFTKWLVKVTKQHNDKCKH
						LLSLMGIPYLDAPIEAEASCAAL
						VKAGKVYPVATEDMDCFTFGS
		l				PVLMQHLTASSDYCKSIWSIGP
						KRAVDLIQKHKSIEEIVQRLDPN
1						KYPLPENRLHKEAYQLFLEPEV
						LGPESVELKWSEPNEEELVKFV
						CGEKQFSEERIHSGVKRLRTEK
				•		GGGEPRGP/DWTQPGRASPTKR
						PDHQEDQHTRSRSSEGSH*GWL
19537	49905	A	19652	1040	2322	LPACPPASTPASTMSIMVTQKS
	•					YKVST/SRWSSGPQAFGNCSYA
		ŀ				RGPGACISSLSFSRVGSSIFRGVL
		ŀ				GGSYGGASGMGSITAVTVNQS
						LLSPLNLEVEPNIQAVCTQEK\T
			:			LNNKFASSIDKVQFLEQQNKML
						ET/ESYINSLKQQLETLGLERLK
						LEAELGNMPGLVEDFKNKYED
						EINKRTEMENEFVLIRQYVDEA
						YMNKVELESCLEGLTEEINLLG
						QLYEEEIQKLQSQILDTSVVLC
						MNNSHSLDMDSIIAEVKAQYEE
						MANHSQAEAESMYQIKYEELQ
						TLAGKHGDDLRHAKTEI/S/EM
						NQNGAKQDMA/*QLHEYQELT
						NIRLALDIKITIYRKLLEGEESQL
						ESGMQNMNIHMKTTSGYAGGL
						SSAYEGLTSPSLSYGLGSSFGSG
						MGSISFSHTSFTRAIVVKKIEIHD
						GKLVSESSDVLPK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19538	49906	A	19653	1040	2322	LPACPPASTPASTMSIMVTQKS
						YKVST/SRWSSGPQAFGNCSYA
						RGPGACISSLSFSRVGSSIFRGVL
						GGSYGGASGMGSITAVTVNQS
						LLSPLNLEVEPNIQAVCTQEKIT
						LNNKFASSIDKVQFLEQQNKML
						ET/ESYINSLKQQLETLGLERLK
						LEAELGNMPGLVEDFKNKYED
						EINKRTEMENEFVLIRQYVDEA
						YMNKVELESCLEGLTEEINLLG
						QLYEEEIQKLQSQILDTSVVLC
						MNNSHSLDMDSIIAEVKAQYEE
						MANHSQAEAESMYQIKYEELQ
						TLAGKHGDDLRHAKTEI/S/EM
						NQNGAKQDMA/*QLHEYQELT
						NIRLALDIKITIYRKLLEGEESQL
						ESGMQNMNIHMKTTSGYAGGL
į	,					SSAYEGLTSPSLSYGLGSSFGSG
					İ	MGSISFSHTSFTRAIVVKKIEIHD
						GKLVSESSDVLPK
19539	49907	Α	19654	152	207	
19540	49908	В	19655	75	402	
19541	49909	В	19656	1	2319	
19542	49910	В	19657	1	1341	
19543	49911	Α	19658	198	1118	LHAPNSLHHHLPRDLL*LLPLPL
						QHRVSDPGLHSALRPMEEHRA
						QS*QPSAPEDAVQA*WGHGGH
-						SPGPDRAGCHHCRGDQERVST
						HHVLPVCHHPDDAYGGCGAG
						WNPDL/PGQMRSHWMSPKIRPA
						NWTRTSWWALPQAPGLSPGAQ
						SWPSFVPRTTPTTPGTTCPTPSW
						*SWRSTSRTSSSLNPFTESLKNS
						LRTSKPFGWSQSAMATPCPLLP
						PASRVELWPETW\PLGQGHATS
						SQWKCVPERKL*QGGEAGGEQ
						LGREPKPSPPSSFLTGQRQEKSP
						EEYCSLLVPLQYFGSRGTHADL
						LHG

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19544	49912	A	19659	l	2707	MPGPHPASASCPPRLFTPRRRAF CPLRICAPLAAAAFCPPRFLPHT RRRAFLLLCRRGFLPPPCLGGG LAAAFFPHPPLPFLRPHRRGFLP HRRRGFLLPPRLGFLPAAAFYPP PPRVFARRGFVPPSPPRLFARRA FCHTPPAAAFSPSAAAAFCPTA DTAFCTLAALAFCRRGFLPPRC RGFLPLRHRRSFLPHRRRGFLPT AAFCHRGFLLATAFYPPRLFAP RCHGFLPPCRSDFFAAAAFCPR RRGFLPLSPPLFAPVTAAFCHRDFL PPPRIFVPAATALRAGAADSA ASSTGVLARAALRGPCGLHRV ALPVEAMGEKKEGGGGGDAA ATEGGTGAAASRALQQCGQLQ KLIVIFIGSLCGLCTKCAVSNDL TQQEIQTPEPAKPAKPASQTRQ PSQPSHPASQASQTSQPAKQPA SQASQPAKPAKPPSQPSQPSHR ASQASQATQPERPASQPASEPS QPSLPASQASQDRQPAKPAKPD SHPRQPSQP/YQPSQPASQASQA SQPAKPSRQPSQPRQPSQLASQ ASQSTQPLKPAKSPVQQSQPSQ PASPDSQASQASQATQPAKPAK PPSQAAQPAKPAKPPSQPSQSSPP
						PSQRAKPAKPARQPSQPSQSS ASQPSQACQPSQPGSQASQAS AAKPGMPAKPARQPSQPSQPA QARHASQAIQPAKPAG*PSQT

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19545	49913	A	19660	1223	2286	RQCFKATEQPPAIHHPSSQPSRQ PSSQPPIYPSPICGRKSGIQPKFQ LNG*MVNG*MDE\SRERHEWP CNYQVRPKAGEKQPGAEANRE TRPEGGTP/RGERGRQGPATRR QKSTPEAGPSDLEKQIARQEAP AGTDEIERQQARPH*AP*KIPVE WVNGEWLNGRMPGDPHEWQC NYQPAIQPAIQPANQLSIHNLW TQFRHSTKIPVEWVNGEWLNG RMPGKHHEWQCNYQPAIQPAI QPANQLSIHNLWTQFRHSTKIP VEWVNGEWLNGRMPEDHHEW QCNYQPAIQPAIQPASQLSIHN MWTQFRHSGQIPVEWVNGSRL HGRQQGIFMHGNATSSHSPSIQ PAIQPAIQPANQLSIHNLWTQFR HSTKIPVEWVNGEWLNGRMQR NIMNGNATISHSPSIQPAIQPAIQ PVNHLSIHNLWTQFRHSTKIPV EWVNGEWLNGRMPGKNHEW QCNYQPAIQPAIQPANQLSIHNL WTQFRHSTKIPVEWVNGEWLN GRMPGDHHEWQCNYQIAPML QSNRTAASHSPSIQPAIQAAIQS ATHLSIPNLWTQVRHSTKIPVE WVNGEWLNGRMPGRDMNGH ATIRYGPKPEKNNRAPRQTEKQ GQKGAHQRREGKARAGNEETE KYSRSRAIRPRKTDSASRSSRGH RRNRKAASAATLGPIEDTNQNK
19546	49914	A	19661	39	599	LHVRDTGLEVKAISIQGPGTLM GASPNQGQNRPHARQRAPPQ/G /PPGEPERRAPLPSGHGEPCRHR PPPFPQPP/AGTQKPLLQGPGGG *PAENAPTAALGSPAPPRGCQA APPPRSGAGRPDLPTLAGPRPAP A\PPPSAAPPPPSGAPSR/PAAG RQRLSGVSSGPSLGWW*VGRG RGLPAFAQIAGHQVGPRRRRTP AGRKPRSPAGPR
19547	49915	A	19662	39	599	GASPNQGQNRPHARQRAPPQ/G /PPGEPERRAPLPSGHGEPCRHR PPPFPQPP/AGTQKPLLQGPGGG *PAENAPTAALGSPAPPRGCQA APPPRSGAGRPDLPTLAGPRPAP A\PPPSAAPPPPPSGAPSR/PAAG RQRLSGVSSGPSLGWW*VGRG RGLPAFAQIAGHQVGPRRRRTP AGRKPRSPAGPR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19548	49916	A	19663	1	379	YFPHTAIRYPPHLNRQDPLKDL
						VSLACDPASQQPGPLNGSGQLK
						MPSHCLSAQMLAPPPPGLPRLA
						LPPATKAATTSEGGATSPTSP/S
						PGIWDSKGLLPSPLLHRPRNPR
						GQHSRPPAHVFGGKLE
19549	49917	Α	19664	3	285	
19550	49918	Α	19665	2	1556	RLEFPGRRFRSAPAPAGPAPPA
						APGMYSSPLCLTQDEFHPFIEAL
						LPHVRAFAYTWFNLQARKRKY
İ						FKKHEKRMSKDEERAVKDELL
						GEKPEVKQKWASRLLAKLRKD
]		ļ		IRPECREDFVLSITGKKAPGCVL
						SNPDQKGKMRRIDCLRQADKV
						WRLDLVMVILFKGIPLESTDGE
						RLVKAAQCGHPVLCVQPHHIG
						VAVKELDLYLAYFVRERDAEQ
						SGSPRTGMGSDQEDSKPITLVT
	ļ					TDFQESFVTSGVFSVTELIQVSR
						TPVVTGTGPNFSLGELQGHLAY
						DLNPASTGLRRTLPSTSSSGSKR
		l]		HKSGSMEEDVDTSPGGDYYTS
						PSSPTSSSRNWTEDMEGGISSPV
						KKTEMDKSPFNSPSPQDSPRLSS
1						FTQHHRPVIAVHSGIARSPHPSS
						ALHFPTTSILPQTASTYFPHTAIR
		İ				YPPHLNPQDPLKDLVSLACDPA
						SQQPGPSWYLG*QRSSSLAPSPS
						SQESQGAAQPAPGPRFRWKIRV
						NKNTPADSQPGQKDKTHRRTH
						SGGKEKTKAE
19551	49919	Α	19666	1	477	
19552	49920	Α	19667	1	440	WTQDM/EGGISS/PVKKTEMAK
						SPFNSPSPQDSPRLSSFTQHHRP
						VIAVHSGIARSPHPSSALHFPTT
						SILP\QTASTYFPHTAIR\YPPHL
						NPQDPLK\DLVS\LACDP\SSQQP
						GP/SYSPPDTSPANR\SFVGLGPR
				İ		DPAGIYQAQSWYLG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence	ŀ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19553	49921	Α	19668	2	1963	CPTGFEHGSPCARRTPTGRVSV
						TRCRRRAWGSLRVSVCGPRAQ
						LQAELVQEEQGGGVLCVIVMP
						LTPGRTPRLPRGRALCNGRGGV
						WGGGGSPRGLLPAQAPGDPEV
						CKFRFLDRCFPPYAGPSDLSLR
						GRAVGPPSASLALLTSRLSGLQ
						DEFHPFIEALLPHVRAFAYTWF
						NLQARKRKYFKKHEKRMSKDE
						ERAVKDELLGEKPEVKQKWAS
						RLLAKLRKDIRPECREDFVLSIT
		ŀ				GKKAPGCVLSNPDQKGKMRRI
						D\CFRQADKVWRLDLVMVILF
						KGIPLESTDGERLVKAAQCGHP
						VL/CALKPQHI\GVAVK/ENLDL
				:		YLAYFVRERDA*QSGSPRTGM
						GSDQEDSKPITLDTTDFQESFVT
						SGVFSVTELIQVSRTPVVTGTGP
						NFSLGELHAHLAYDLKPASAGL
						TRTLSSTFSSGNKRY\KAGSMEE
						YVDSTPWGDD*T*PSSPTSSSRN
						WTEDMEGGISSPVKKTEMDKS
						PFNSPSPQDSPRLSSFTQHHRPVI
						AVHSGIARSPHPSSALHFPT\TSI
						LTPDGPSPTFPHTAIRYPPHLNP
						QDPLKDLVSLACDPA\SQQP\GP
						LNGSGQLKMPSHCLSAQMLAP
						PPPG\LPRLALPPATKPATTSEG
						GATSPTSPSYSPPDTSPANRSFV
						GLGPRDPAGIYQAQSWYLG
19554	49922	A	19669	332	551	ESWLGRGHQPPP\PGLFHSPKPS
.,,,,,,						SGTAVLPAWRPPSPGPGAGDD
						ASPSPS*AGPGAGANPGA*QRS
						ASGEKGS
19555	49923	Α	19670	1614	1673	
19556	49924	Α	19671	72	227	EEFPMGHTA*WSCEKRATILQI
						QEW*IQQQLAPCAWQSHRHST
						RAHESSHG
19557	49925	С	19672	1	604	
19558	49926	Α	19673	653	889	VYPQTNWKQNTGTRVFIAALF
						TM\AKGENK*PSMGEW/IKQM
						WSIHTMERHAAIKRIEAQVYAT
						M*MKLKSMLLLERNQT

SEQ ID	-		SEQ ID NO:	L		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/340,217	sequence	or peptide sequence	deterion, (-possible nucleotide insertion)
					•	
19559	49927	Α	19674	75	1619	EVSIPAQTMWTADEIAQLCYEH
						YG\IRLPKKGKPEPSHEWTLLA
						AVVKIQSPA\DKACDTPDKPVQ
						VTKEVVSMGTGTKCIGQSK/M*
						RKNGDIL\NDSHA\EVIA\RRSFQ
						RYLLHQLQLAATLKEDSIFVPG
						TQKGVWKLRRD\LINVFFSSHT
						PCGDASIIPMLEFEDQPCCPVFR
						NWANNSSVEASSNLEAPGNER
						KCEDPDSPVTKKMRLEPGTAA
						REVTNGAAHHQSFGKQKSGPIS
						PGIHSCDLTVEGLATVTRIAPGS
						AKVTDVYRTGAKCVPGEAGDS
						GKPGAAFHQVG\LLRVKPGRG
						DRTRSMSCSDKMAR\WTVLGC
						QGALLMHLLEEPIYLSAVVIGK
						CPYSQEAMQ\RA\LIGRCQKV\S
	i					ALPKGFGVQELKILQSDLLFEQ
						SRSAVQAKRADSPGRLVPCG\A
1						AISWSAVPEQPLDVTANGFPQG
					,	TTKKTIGSLQARSQISKVELFRS
İ						FQKLLSRIARDKWPHSLRVQKL
						DTYQEYKEAASSYQEAWSTLR
L		L				KQVFGSWIRNPPDYHQFK
19560	49928	C	19675	45	140	+ WW. TO WOOD WOOD AD TO SO
19561	49929	Α	19676	3	354	AKIKTGYK\REHINLGCDMDFDI
						VGPSIRGALVLGYEGWLAGYQ
						MNFETAKSRVTQSNFAVGYKT
						DEFQLHTNVLLLNLERTCFLTG
						PTPTPPPLSWLPGTTFSLTWAM TGRYTINSQ
19562	49930	A	19677	 	983	AARSAPLPGSAPCLRVAAPAVA
19302	49930	Α.	19077	1 1	763	ASEPAATSSEQKMAVPPTYADL
						GKSARDVFTKGYGFGLIKLDLK
]					TKSENGLEFTSSGSANTETTKV
						TGSLETKYRWTEYG\LTFTEKW
						NTDNTLGT\EITV\EDQL\ARGT*
1						SLTF\DSSFSP\NT\GKKNAKIKT
						GYKREHINLGCDMDFDIAGPSI
						RGALVLGYEGWLAGYQMNF*
						DCKIPE*PQSNFWQLANKT\DEF
						QASR*M*ND\GTEFGGSIYQKV
						NKK\LETAVNLPWT\AGNSNTR
						FEIAAKYQ\VDP\DACFSGKVNN
						SSLIGLGYT\QTLKPGIKLTLSAL
						\LDGKN\VNA\GGHKL\GLGLEF
						QA
L	<u> </u>	<u> </u>	<u> </u>	L	l	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19563	49931	A	19678	551	831	CHPSSHVLCAPASLSDHQWPPP ALCPSSRPQ/PCALPWSGTLSSR PQPCPTHPSGIEA*LCLPLVTQL G*LPL\PAPQHLVPPCPSTCPTM LKLF
19564	49932	A	19679	47	980	VCQYCTARMADFGISAGQFVA VVWDKSSPVEALKGLVDKLQA LTGNEGRVSVENIKQLLQCLVP GSTT\LHSA\EILA\EIARILRPGG CLFLKEPVETAVDNNSKVKTAS KL\CSALTLS\GLVEEKELQREP LTPEEVQSVREHLGHESDNLLF VQITGKKPNFEVGSSMQLKLSI TKKSSPSVKPAVDPAAAKLWT LSANDMEDDSMDLIDSDELLDP EDLKKPDPAS\LRA\ASCGEGKK RKACKNCTCGLAEEL\EKEKSR EQMSSQPKSA\CGNCYRGAMPS GCASCPYLGMPAFKPGEKVLLS DSNLHDA
19565	49933	A	19680	29	1270	FPFWPAVFQVCQYCTARMADF GISAGQFVAVVWDKSSPV\EAL KGL\VDKLQAF/TPGNEGRVSV\ ENIKAAVAILPTKNPSFGHYFV Q/CLVPGKAPLWHSA*DFWAG NPPGFLRPGWMFFFLKEPVETA VR*Q*AKWKTASKL\CSAL\TLS GLV/EKLKELQREPLTPEEVQSV REHLGHESDNLLFVQITGKKPN FEVGSSRQLKLSITKKSSPSVKP AVDPAAAKL\WTLSANDMEDD SMCIFCGCSLTHRWPLEHVVQV E/IMMDQPKRRTRVDT\FFTPRT PKFPSRSPASHFSFSIKQKT/TRP VSLIALNTL\QDLIDSDELLDPE DLKKPDPSLLRAASCGEGKKR KACKNCTCGLAEELEKEKSRE QMSSQPKSACGNCYLGDAFRC ASCPYLGMPAFKPG\EKVLLSD
19566	49934	A	19681	3	405	QDFGTRALAAVQIKNGGSWWC PPVSLAAGVGPRPQLGRPYVRV APHGQV\GLSFAMWKTLT\FFR GASPGVGSQAMLKCVA*SPHH WRARRDPEFIAYP\HLR\IRTKP\ FPWG\DGNHTQQYITPH\VNPLS NLATED

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	•	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19567	49935	A	19682	2	450	VNKAGGLIYQLDSYAP/RAEAE
						KTFSYPLDLLLKLHDERVLVAF
						GQRDGIRVGHAVLAINGMDVN
						GRYTADGKEVLEYLGNPANYP
						VSIRFGRPRLTSNEKLMLASMF
						HS\IKFVVLADP\RQAGIDSLLRK
						IYEIYSDFALKNPFYSLEMP
19568	49936	Α	19683	1	2034	
19569	49937	Α	19684	202	453	RRLWRGSCPRCGGKARDGMG
						VRGR/YSEG/RERV*FGGPPRGS
						TWPLCGRWRRPARPSAPLRWS
			<u> </u>			AMCPNTSDAAGACAAAVRCG
19570	49938	Α	19685	269	326	
19571	49939	В	19686	98	562	
19572	49940	Α	19687	1	2223	
19573	49941	Α	19688	374	1107	IPVSERQGSDWRFFSVYVVNKA
						GGLIYQLDSYAPTGLRLEKTF\S
						YPLDLAGSS*HDERVLVAFGQR
						GRHPEWGHAVL\AINGM\DVNG
			}			RYTADG\KEVLEYLG*PLLNYP
						VSIRF\GRPRL\TSNE\KLMLGLP
			,			CFHSLFAIG\SQACLPGTREASR
			ļ			HLKILGRQDTFK\LH\CYQTLT\G
-						IKFVVL\ADPR\QAGI\DSLLRKI
						YE\IY\SDFALKNP\F\YSLEMPIR
			İ			CELFDQNLKLALEVAEKAGTF
19574	49942	Α	19689	119	288	ALHLNSSDVHQKLLPVQSLKKS
						*GRHRNNTCNQRVHA/SSTSRE
						FFTKLNKDNDILKP
19575	49943	Α	19690	92	317	KSVGPVNFPLPSGNMNEDN*M
						H*HPFSSVTQAFRLRVLQSII\QR
						PL*TFNPGSSSLFSSISGQSNFPL
						GVAVLRP
19576	49944	Α	19691	27	389	NFKTLKNN*IYK*MCSQEIKILS
						*KLIRFWYEIHAFFPWAHGVVF
		1				CVAQEFCLPTVVLLPVPRLLIM
		1				KDMVRRLQELRHTEQVQRAYA
						LNCGEGATVSYEIQIRVLREFGL
						ADAAAELLQN
			I			ADAAAELLQN

19577 49945 A 19692 313 2255 KTPQVQPFPRQTGAAGAGAP SGRSGERAAGRAGPSGGSGGG QDSAHAGEAYRVRGILLRKTFQ CLKDYFLTYYTSSKVVPVKVPC RQVLTIDKCSLPRSSHRNARTV SNDFPRKSCSRERCTHCMRPPE GRRPPNGMNSWKWPSGDESK WRRGEPGAVPEPEPLTGEERAA AAAVGGGWPGVLAPVRTAVL VAAAAEATAAVPAAAAAAGG ASGHCNHPVMQGRNLSHRFLL PLHHSPYQFGTSSYSQQYGCE SKLYSLDHGHEKPQDKKKRTS GLATLKKKFIKRRKSNRSADHA KQMRELLSGWDVRDVNALVE EYEGTSALKELSLQASLARPEA RTLQKDMADLYEYKYCTDVDL IFQETCFPVHRAILAARCPFFK/I TAFFLTRVWGRDNNGHQYSW Y*YAHVFCFVTLPLYRRVWEW EDSRFQNVDILVQLSEGIWEHQI PLM*DMRGTLLI/DMCYYDVVL SFSSDSELVEAFGGNQNCLDEE LKAHKAVISARSPFFRNLQRRI RTGEEITDRTLRTPTRIILDESIIP KKYATVILHCMYTDVVDLSVL HCSPSVGSLSEVQALVAGKPN MTRAEEAMELYHIALFLEFNM LAQGSCWECIPPNDEANQEQRS EGSRKWAPTREGSKGRSHASA	SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
EGSRKWAPTREGSKGRSHASA	19577	49945	A	19692	313	2255	SGRSGERAAGRAGPSGSGGG QDSAHAGEAYRVRGILLRKTFQ CLKDYFLTYYTSSKVVPVKVPC RQVLTIDKCSLPRSSHRNARTV SNDFPRKSCSRERCTHCMRPPE GRRPPNGNMNSWKWPSGDESK WRRGEPGAVPEPEPLTGEERAA AAAVGGGWPGVLAPVRTAVL VAAAAEATAAVPAAAAAAGG ASGHCNHPVMQGRNLSHRFLL PLHHSPYQFGTSSYSQQGYGCE SKLYSLDHGHEKPQDKKKRTS GLATLKKKFIKRRKSNRSADHA KQMRELLSGWDVRDVNALVE EYEGTSALKELSLQASLARPEA RTLQKDMADLYEYKYCTDVDL IFQETCFPVHRAILAARCPFFK/I TAFFLTRVWGRDNNGHQYSW Y*YAHVFCFVTLPLYRRVWEW EDSRFQNVDILVQLSEGIWEHQI PLM*DMRGTLLI/DMCYYDVVL SFSSDSELVEAFGGNQNCLDEE LKAHKAVISARSPFFRNLLQRRI RTGEEITDRTLRTPTRIILDESIIP KKYATVILHCMYTDVVDLSVL HCSPSVGSLSEVQALVAGKPN MTRAEEAMELYHIALFLEFNM
11057X 1/100/16	19578	49946	В	19693	4	3923	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19579	49947	A	19694	1	2154	MEYYAAMKTKEIMSFALTWIE
1,3,7	',	ļ ` `	1203.	1		LEAIVLCELTHVLSELMQKQKT
						KYHMLSLGSPVPTFALWNFDV
						AQDFHHFIENARSSQCRGRPRM
						TGAGAGAPSGRSGERAAGRAG
						PSGGSGGGQDSAHAGEAYRVR
			:			•
		ŀ				GILLRKTFQCLKDYFLTYYTSS
İ						KVVPVKVPCRQVLTIDKCSLPR
		İ				SSHRNARTVSNDFPRKSCSRER
ŀ						CTHCMRPPEGRRPPNGNMNSW
İ						KWPSGDESKWRRGEPGAVPEP
ŀ						EPLTGEERAAAAAVGGGWPGV
-						LAPVRTAVLVAAAAEATAAVP
						AAAAAAGGASGHCNHPVMQG
						RNLSHRFLLPLHHSPYQFGTSS
						YSQQGYGCESKLYSLDHGHEK
•						PQDKKKRTSGLATLKKKFIKRR
•						KSNRSADHAKQMRELLSGWD
İ						VRDVNALVEEYEGTSALKELSL
						QASLARPEARTLQKDMADLYE
						YKYCTDVDLIFQETCFPVHRAI
ŀ						LAARCPFFK/ITAFFLTRVWGRD
İ						NNGHQYSWY*YAHVFCFVTLP
						LYRRVW\KEDSRFQNVDILVQL
						SEEFGTPNSLDVDMRGLFDYM
						CYYDVVLSFSSDSELVEAFGGN
						QNCLDEELKAHKAVISARSPFF
						RNLLQRRIRTGEEITDRTLRTPT
						RIILDESIIPKKYATVILHCMYTD
		ŀ				VVDLSVLHCSPSVGSLSEVQAL
l						VAGKPNMTRAEEAMELYHIAL
					,	FLEFNMLAQGSCWECIPPNDEA
19580	49948	Α	19695	1	163	
19581	49949	Α	19696	1	207	
19582	49950	Α	19697	1	579	PTRPGQAGSSSAMAAQRLAKR
						VL\SKLQSPSRARG/PCAGSPGG
						LQK/RGHARVTVKYDRPGSLQR
						RLDVEKWIDGPLEGLYRR\QEA
						D\MPMRSNIDELLE\LESEE\ERS
						RKIQGV/LRDS*KSCGKPVEDFI\
						QELLAKLQGLHRQ\PGLRQPSPS
						HDGSLSPLQ\DRARTASPLTLLA
						LFPGPPERRPALLCVLSCI
19583	49951	A	19698	2	151	PPSPPPLPPPPLPPAPSLPLSLPSF
						DLPQPPVLDTLGV/LF*LCRKKI
						SVS
19584	49952	Α	19699	2	310	
		لستت		<u> </u>		

NO:	of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19585	49953	A	19700		688	LTWMTERVDFTALLPLEPPLPP GTLPQPSPTPPDLEAMASLLKK ELEQMEDFFLDAPLLPPPSPPPL PPPPLPPAPSLPLSLPSFDLPQPP VLDTLDLLAIYCRNEAGQE\EV G\MPPLPPPQQAPSFLLPP\QPSR WGPLTHIPGQPTRGGPAKQKEE RPRNKSAALKVTPSGKRARG*S PWRAKSQGLEGTEFASLKERA\ ESVEREIQ\YVKD\LLIEVYKARS
19586	49954	A	19701	3	902	QRTRSC AAMATPARPGEAEDAAERPLQ DEPAAAAAGPGKGRFLVRICFQ GDEGACPTRDFVVGALILRSIG MDPSDIYAVIQIPGSREFDVSFR SAEKLALFLRVYEEKREQEDC WETFGGAGGEASPA*RRSSSS GTRRWTWRTL*LGSSATATCW PC\GESDRQVLDLD/LGSTNARS SCARGRAGSGTCQGPSSWGPR G/CYSWYKGQPKTCFKCGSRTH MSGSCTQDRCFRCGEEGHLSPY CRKGIVCNLCGKRGHAFAQCP KAVHNSVAAQLTGVAGHLNTR LPARVNTQASLSLLKCPKLFF
19587	49956	A	19702 19703	1359	1981	QSPFNDTLPGVPVSA*LHSLPPE HHPSKG*GVAKSPPVYPAGILL VCNNCAAYRKLLEAQTPSVRK WALRRQNEPLEVRLQRLERER TA\KKSRRDNETPEEREETDEQ RARRLQRDREAMRLKRANETP EKRQARLIREREAKRLKRRLEK MD\MMLRAQFGQDPSAMAAL AAEMNFFQLPV\NGVEL\DSQLL GKMAFE\EQNSSSLH

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19589	49957	A	19704	1	673	MSAKDGTCTLQDCMYKDVQK
						DWPGYSEGDQQLLKRVLVRAH
		1				NTQRQRQNERSDTRPCQRAAP
						RRFSHGTRRGTLPLASPDRARA
	1	ł				PRTGHCRTQRRLDRTGGGFPPL
						TGAWAMLADWIHEGTPSRGRS
						DNDEPSAGSVGGLRQHGPGPR
						APKLTTTKVKALASPVLGQKIA
						ALCLGLTAQGLNIMDNPGSLAI
						HVHPDSVLTRKTNPCTHIWGFT
						QREPPLVANMSAKDGTCTLQD
						CMYKDVQKDWPGYSEGDQQL
						LKRVLVRAHNTQRQRQNERSD
						TRPCQRAAPRRFSHGTRRGTLP
						LASPDRARAPRTGHCRTQRRLD
		ļ				RTGGGFPPLTGAWAMLADWIH
						EGTPSRGRSDNDEPSAGSVGGL
		1				R/HAWPRTKSPKANNH*SESTG
						QPGAGPKNCSPVLGAYSPGPQH
						HGQPGLPGHTCP/SRTQC*PGRP
						TPAPTSGVSLRGNPLWWPT
19590	49958	Α	19705	1088	2273	SPCLLTGGHLPADTSYRRAPAG
		İ		:		IWQVPLWDEASQRKE*YQHQQ
				İ		KKMSTQKPHPKDHNSSPAREQ/
						SSMENKFHELTEVGFRKITSLE
						KNINDLMELKNTARELREAYTS
						IKSRIDQAKESISEIEDQLNEIKC
						EDKVREKRMKSNKQNFQEIWD
						YVKRPNLRLTGVPESDRENGTK
			1			LENTLQEIIQENFPNLARQANIQ
						IQEIQRTAQRQSSSRATPIHIIVR
						FAKFEIKEKMLRAAREKDISMR
						QKINKDIQDFNTVLDQVDLIHV
						YRTVHPKSSEYTFFSAPHHTYC
						KTGHIIGSKTLLSKCKRMEIITN
						TLSDYSAIKLELRIKKLTQNHTT
					_	TWKVNTLLVNEY

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	l	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19591	49959	Α	19706	318	873	AYELTPEPSPLPVFPVSADGSTT
						HPLADKRKASPVPEQHEEHLPA
						RPVTQRCLTYPFPNITTASEFGP
						ATGFRARRSGRGSTVCCNPGCY
						PRARPEPEPRGTPPSVSAPRAAQ
						EAPARQR\AGPVLADTVP/SPAP
						RGKRDISPLFGNAHSASPKPSLD
						LWAPPP*AAGRRGPQQA*DVTE
						DTQQQWSAMAVGGTAKWM
						AALGRFLVTYVEVEGVKLQTF
	:					TVTTLTQSFKYSLSKYLQQQPL
						LSSRTIYPTFYYLSLLRNLRAFK
						LNAAEAKPMSSPQNLVLFLCFL
						SLQMAAPPIRLQTKGKQAQCQS
						STKSTFQRGRSLRGVSPTLSPTS
						QQPRNLDQQLASEQGEVGEAP
		1				LCAVIQGVIPVLGRSRSREEHPP
		1				VSALPGLHRKRQRDKEAGPVL
						ADTVPFRPHAGKGTFLRCLETL
						TRRAPNPRWTFGLPLRKLQGA
						VAHSRPEIRTHSVPNRT
19592	49960	Α	19707	1	690	DQDMTSSDIPVRLEGSLWLP*IF
						IGTFKAFDKHMNLILCDCDEFR
						KIK*GVDCGGRRLEDGNGLSG
						WAKQ*SLVSMTVEGPPPKDVS
		İ				QSRKP/CWRRGGPTVSQALC*IL
						YPSWGSHAPGSCRTCWASPWG
						WRAIPTGEELGKGYHLLVAKIK
						DKGEGP*FSNPVPTWPWGSSPT
1						YGPRSTPSR*GAHKETSAN**K
				}		MP\SWINDEI*PPMGMPPPGMRP
						PPPGMRGKCLQW

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:		Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ĺ	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19593	49961	Α	19708	1	1348	MVFWARPRTLLLCAALGLGAL
						YTATPAPTVAKRGQGTAQALS
						SEGASPKPWWLRHGIEPCKQSY
						WEVRTEQSPGVSERKAAAPVR
						GLEIKLPSPWDKAPEGGGGCGH
ĺ						SFSRLKCSFMPVLKRAVDLPAQ
						RSSPANGQTACSSGSLAPVLPD
						WEIPPSKGRQTPYATGELRLAS
						DGCPSGTKLPEEGTDSNLCCSA
						SSAESNNININININKKDAHGKT
						PSKGPQHQRSKLDKSMKTRKN
						HHKNAENSKNLNASSPPKDHN
						SSTAKEQNWTENEFDELTEVGF
						RRRVIANSSKLKEHVLTQCKEA
		•	<u> </u>		İ	ENFEKILEELLTRITSLEKNINDL
			<u> </u>			MELKNTAQEHREAYTSINSQID
						QAEERISEIEDKLNEIKREDKIR
		İ				GKRIKRNKK\SLQEIWDYVKRA
						NL*LIAVVPLNKITSPRTSETLST
		1				STNMSRSTLSCLLSSFSCPASCT
						ATSAALF
19594	49962	Α	19709	718	1138	GGRAAEGSQPPVAEPRCSRGAR
						ALPLPDTSWRQPS/APQAAP*D
						VELPASPSAPFAVLLSKPRLSQN
}						KPKSPFSVASP*HGDKRPQPSRH
						LI*IPHLSSRAPVHS/PLGWPSNR
			1	1		K/HKHLRPDTAHAPAEGPPLNY
10.50.5	100.50			<u> </u>	7.00	VRKQQKQS
19595	49963	Α	19710	5	760	RAAACGGAAPGRKRCAPGRKR
						CGPRRKRSLGPPTLGQRPRQPS
			1			QQLRKPRRRRGSPGSLPKSSPLP
						SSTGKTGWSTRGRSVRPPVTTS
						RCCPPGHPCSRRAPSCTARAPLP PT*PPTTAPCRASPPAPRPPRRPR
]			PPPGTRSPTSRGRCRWTRSGAG
						TSRCSWRAGPGTAPAAAGARS
		ŀ				RPGAPGVSAAVPCAAACSWMT
						SAPEVRAASSPGTPSRGSGGRS
						HPPCSRGLPTTGPATTAPEAPG
						CWSCRRS
19596	49964	В	19711	180	300	o John de
19597	49965	A	19712	249	700	RQDPVSGGQSCPASCPSLWAA
			-			ARVWLGQ\PAPRAR*CPVCSSC
						PGPGAPEDTPEAGGCGCSGCRC
						CLRTQEPPAPLQPPHPPMAQTP
						REEVGRRGWTPGCPGTDPCLW
			[GLRLLCPLWDPHRSGTRPWAQ
						VQRDTTETKAQRGLVRCQRGC
19598	49966	С	19713	273	485	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19599	49967	Α	19714	35	779	VRPPSHVTADSGRSPLSLTYLPL
		1				QEPGDMAAAVPRAAFLSPLLPL
		1				LLGFLLLSAPHGGSGLHTKGAL
		1				PLDTVTFY/KDYGDKLNMELSE
						KYKLDKESYPVFYLFRDGDFEN
		1				PVPYTGAVKVGAIQRWLKGQG
		1				V\YLGMPGCLPVYDALAGEFIR
						ASGVEARQALLKQGQDNLSSV
		1				KETQKKWAEQYLKIMGKILDQ
		1				GEDFPASEMTRIARLIEKNKRSC
		1				KKAVCDFPGFGGGREGRVNLL
		ļ <u>.</u>				AVSPLWNIRG
19600	49968	Α	19715	2	253	
19601	49969	Α	19716	494	1443	LAVPRPAYGPRSPGSLLRIPVPG
		İ		; ;		LHLLNLNVWSRGSAAALHTKG
						ALPLDTVTFYKVIPKSKFVLVK
	-	İ				FDTQYPYGEKQDEFKRFAENSA
		ŧ	ŀ			SS\DDLL\VAEVGISDYGDKLNM
		ŀ	<u> </u>			\ELS\EKYKL\DKESYPVF\YLFR
						DGDF\EEPSSHYTGGS*RLGAIQ
						RWŁKGGKGVYLGDGLELVCT V*LTPLAGEF\IR\ASGVEARQAP
						LEAGAQDNLSKCERRLRKKW
	1					AEQYLKIMGKILD\QGEDFP\AS
						EMTTESLGLIEKN\KMSDGKKE
	1					EL\QKSLN\ILTALPRRKGAEKR
						GAVNKAVCDFPRVWWEYGRG
						ELTLLAVSPLWNIRG
19602	49970	A	19717	286	332	ННННРНҮННННРНҮННН
17002	1.33.0	1				HPQYHH\HHPHYDHHRHRQGD
						HHGQLQYDDDHQPQCHQHDH
						LRCEHQHYIQ*WLINNLRASRY
						KPLLNIMLVFAAEMIVLVALGL
						MVVVVLELTVVVALTMTVVVI
						VRMMGGDIGDDGGGGSGDDG
						GGGSGDDGGGVMWQFFDSMS
			-			KNDSIIPTTTTTIIPTTTTTTIRA
						EQFTILTWKPVVSSQKNTSSFR
19603	49971	С	19718	.1	546	
19604	49972	Α	19719	1	90	
19605	49973	В	19720	217	951	
19606	49974	В	19721	67	1708	
19607	49975	В	19722	218	3267	

SEQ ID NO:	of peptide	Met hod	SEQ ID NO:	location of first	codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19608	49976	A	19723	1	634	LGRLSFPRPEQPDPGHRAHPSSP PRPRRPHARGRRGPRRHH/YHD
		1				TGANRAPEVRPPCPES/SPVGVA
						AFEPNGLRGQQPSPRSPGAPKG
						L*PTPPSASSSPKPREPSALGRPC
						CRCEESRFLEHRNSNPRVKDSR
						QARGRHAEFHGRTGVFERRRS
						RDHLPTHFPASVRWEKLSWCA
						RKNGLSGLRLCHGSQHFSGYSN
						RKVGTESLNQVWE
19609	49977	A	19724	3	1090	
19610 19611	49978 49979	A	19725 19726	23	636	FSYLPPGPSHGTWGLWELQFK
19011	149979	A	19/20	23	030	MRFGVCRHLMEDSMDMDVSP
						LR/PQNYLFSCELKADKDDHFK
						VDNDENEHQLSLRTPTVSLGVF
						EITPPVLLWLKCGSGPVHISGQ
						HLVAVEEDAESEDEEEEDVKLL
		1				RISGKTKTFMATNGKEYKHYKI
		1				SSEKSLDNKYKTRTPGFQAFGF
		1		-		EDLHPWPLGSQAFYLSLRVTPP
		1				VFLVLRLLDFD
19612	49980	Α	19727	1	830	MEDSMDMDMSPLRPQNYLFGC
						ELKADKDYHFKVDNDENEHQL
						SLRTVSLGAGAKDELHIVEAEA
						MNYEGSPIKVTLATLKMSVQPT
						VSLGGFEITPPVVLRLKCGSGPV
						HISGQHLVAVEEDAESEDEEEE
						DVNRVLFMKP/KRQGPQAYVR
						GFGVQLEIGSH*DPEQQF*RDSA
						QPSLATPSTPHLQALLCLSGQV
						LRLPLWGQRL*GLQGLLPPQHP
						EEHGVHVSPGQELHHQQGDPE
						PLPVLPTAEVL*SGHVQGVCEK
19613	49981	A	19728	1	684	RPKQEEEGGAQARVL
19614	49982	A	19729	3	535	DSVLRGCSLEQRSFISVRLLSYL
						SACRHPMEDSMDMDMSPLRPQ
						NYLFGCELKADKDYHFKVDND
						ENEHQLSLRTVSLGAGAKDEL
		1				HIVEAEAMNYEGSPIKVTLATL
						KMSVQPTVSLGGFEITPPVVLR
						LKCGSGPVHISGQHLVVYRRK
						HQELQAMQM\DCRALSTS*ASS
10615	140002	I.	10720	1	2260	APRPS
19615 19616	49983 49984	B A	19730 19731	1198	2268 472	AHDIGCPEVSPLVLEVARDEGP
19010	147704	\\\^	19/31	190	7/2	AHRPRFSH*SSHCSHWGTSQRA
						APPPCHPCIQTTW/PGAVTGLG
						G*PKTRSGPGKDRSSGAASSFP
	1					GGRV
L		Щ	L		I	OGICY

SEQ ID NO:	SEQ ID NO: of peptide sequence	ł .	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19617	49985	A	19732	92	2049	LTALPPAVSPGGHSSSSSSRTIS GIT*ELQR/LLQAVVPGPWQED VADAEECAGRCGPLMDCRAFH YNVSSHGCQLLPWTQHSPHTR LRHSGRCDLFQEKGEWGYMPT LRNGLEENFCRNPDGDPGGPW CHTTDPAVRFQSCGIKSCRVAA CVWCNGEEYRGAVDRTESGRE CQRWDLQHPHQHPFEPGKFLD QGLDDNYCRNPDGSERPWCYT TDPQIEREFCDLPRCGSEAQPRQ EATSVSCFRGKGEGYRGTANTT TAGVPCQRWDAQIPHQHRFTPE KYACKDLRENFCRNPDGSEAP WCFTLRPGMRVGFCYQIRRCT DDVRPQDCYHGAGEQYRGTVS KTRKGVQCQR/WSLE/TG*DYP ATTSLPLPPRFTFTSEPHAQLEE NFC/RDPDGDSHGPWCYTMDP RTPFDYCALRRCDQVQFEKCG KRVDRLDQRCSKLRVAGGHPG NSPWTVSLRNWQGQHFCGGSL VKEQWILTARQCFSSCHMPLTG YEVWLGTLFQNPQHGEPGLQR VPVAKMLCGPSGSQLVLLKLE RYVDNLGGWTKCEIAGRGETK
						GTGNDTVLNVALLNVISNQEC NIKHRGHVRESEMCTEGLLAPV GACEGDYGGPLACFTHNCWVL KGIRIPNRVCARSRWPAVFTRV

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19618	49986	A	19733		2296	HSAGQRSPSNDFQVLRGTELQH LLHAVVPGPWQEDVADAEECA GRCGPLLLSRAFHYNVSSHGCQ LLPWTQHSPHSRLWHSGRCDLF QEKGEWGYMPTLRNGLEENFC RNPDGDPGGPWCHTTDPAVRF QSCGIKSCRVAACVWCNGEEY RGAVDRTESGRECQRWDLQHP HQHPFEPGKFLDQGLDDNYCR NPDGSERPWCYTTDPQIEREFC DLPRCGSEAQPRQEATSVSCFR GKGEGYRGTANTTTAGVPCQR WDAQIPHQHRFTPEKYAC/KVR WAGGRALGRAAAGETGGG*SR A*GWRLAG*G*VERLLRDLREN FCWNLDGSEAPWCFTLRPGTR VGFCYQIRRCTDDVRPQDCYH GAGEQYRG\TV\SKTRKGVQCQ RWSAETPHKLQ*VPGAPGPAR ALTLGRHALMSGTRAWKWLR LSLPRFRSRSRLGSRLTSEPHAQ LEENFC/RDPDGDSHGPWCYTM DPRTPFDYCALRRCADDQPPSIL DPPPDQVQFEKCGKRVDRLDQ RRSKLRVAGGHPGNSPWTVSL RNR*GTTACLPQRGAEVVSSVV MPLGAGNLSLPPEVLARRWQV *HLSQESVPCPNSPLL*\QGQHF CGGSLVKEQWILTARQCFSS\CE PPLCLGTQSHPTFPFPQAS*QVS LGATD*ESQMT*QSFSPSHMPL
						TGYEVWLGTLFQNPQHGEPGL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hođ	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19619	49987	A	19734	727	3012	 PAPTHAHSAGQRSPLNDFQVLR
19019	49967	 ^	19734	1,2,	3012	, , ,
						GTELQHLLHAVVPGPWQEDVA
						DAEECAGRCGPLMDCRAFHYN
]					VSSHGCQLLPWTQHSPHTRLRR
						SGRCDLFQKKDYVRTCIMNNG
						VGYRGTMATTVGGLPCQAWS
						HKFPNDHKYTPTLRNGLEENFC
						RNPDGDPGGPWCYTTDPAVRF
	ŀ					QSCGIKSCREAACVWCNGEEY
						RGAVDRTESGRECQRWDLQHP
						HQHPFEPGKFLDQGLDDNYCR
						NPDGSERPWCYTTDPQIEREFC
						DLPRCGSEAQPRQEATTVSCFR
						GKGEGYRGTANTTTAGVPCQR
						WDAQIPHQHRFTPEKYACKDL
						RENFCRNPDGSEAPWCFTLRPG
						MRAAFCYQIRRCTDDVRPQDC
]		YHGAGEQYRGTVSKTRKGVQC
						QRWSAETPHKPQFTFTSEPHAQ
				Ī		LEENFCRNPDGDSHGPWCYTM
						DPRTPFDYCALRRCADDQPPSIL
						DPP\DQVQFEKCGKRVDRLDQR
i						CSKLRVAGGHPGNSPWTVSLR
						Į.
						NWQGQHF\CGGSLVKEQWILT\
	ŀ					AR\QCFSS\CHMPL\TGY*GMVG
						ATFFQNPQHGEPSL\QRVQ*AK
						MLCGPSGSQL\VLLKLERSVTL
						N/HRVWALIILP\PEWYVVPPGT\
						KCEIAGWGET\KGTG\NDTV\LN
				1		VALLNVISN\QECNIKHRGRVRE
						SEM\CTEG\LLAPVGACESDYG
						GP\LACFT\HNCWV\LEGIIIPN\R
19620	49988	Α	19735	66	234	WCSAWHSWLPHSPSGPAASYP
						PAACRSAPQWPDP*R*CPRTHR
İ						RRSPWRAPWHCP
19621	49989	Α	19736	3	500	GRGGGGVRRCARPAPGRPPAA
						ARRRGDQRGGRLGR\HVPGKTF
						CCGICGRGFGRRETLKRHERIH
						TGEKPHQCPVCGKRFRESFHLS
						KHHVVHTRERPYKCELCGKVF
						GYPQSLTRHRQVHRLQLPCAL
				İ		AGAAGLPSTQGTPGACGPGAS
	L	L			L	GTSAGPTDGLSYACSD

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19622	49990	Α	19737	899	2348	GRRRGGGGRCWRGCQSGPERF
' ' ' ' ' ' ' ' ' ' ' ' ' ' ' ' ' ' '						SCATCGQSFKHSW/DLVTHKYV
						HLMLLLPPSCEAVAPRGWRRSC
						SCPECCRVC/ELSSAA/RSATAR
						CFRRRSPSHAQPAPGPQVPQDW
		}				RSPRARIRLPVGPVGMPAFPTSA
		1				SGKAWPSLRSVSPHAFWKYHT
					1	SPPAMQPSGPLWASVPACAP*K
ŀ		ĺ				NIPSPTIWQRLAFSGLGLPICWV
		ł				ENHGPATSSQWLVSGPRSPHLL
						S/RAEHPFSRPLPRWVATAERQ
		1				RRPSGKRTNPFPASPAPPPGA/G
						PEVSVPPAGALPGEALGPARDL
			ŀ			WPCSSWT*AGLPALG/DTGGGE
						APAP/AAAAEPSEDTLYQCDCG/
						TFFASAAALASHLEAHSGPATY
		•				GCGHCGALYAALAALEEHRRV
		ŀ				SHGEGGGEEAATAAREREPAS
		ŀ				GEPPSGSGRGKKIFGCSECEKLF
						RSPRDLERHVLVHTGEKPFPCL
						ECGKFFRHECYLKRHRLLHGTE
						RPFPCHICGKGFITLSNLSRHLK
						LHRGMD
19623	49991	Α	19738	2	358	
19624	49992	A	19739	635	2441	LKAIKRHPFFVTIDWNVSCMPQ
						SPPPFKPAVGRPEDTFHFDPEFT
						ARTPTGVCHSGSLYNAHHLFR
						GFSFVASSLIQEPSQQDLHKVPV
						HPIVQVTVFACSGFTDGYEIKE
						DIGVGSYSVCKRCVHKATDTE
						YAVKVRQAC*VSSLVFPAGTG
		ŀ				APGIGEYGSGKLVSILCAQDSP
						HNLLPQRKVEKPSLSILVGESRS
						HVDKGQEVDILEMGMEFFFFIE
						FLVSPLATG*CFPMARSPCMGC
						SLSVCTAL*GQPLQGNTPTFKY
		1				QQVTWPQAKPPTPTAVLSPETA
		1				NVTSPQEQTRCLLKDRWLYPV
		1				VRNSSTRGPSEEIEILLRYGQHP
		1				NIITLKDVSGPRATASLVMELM
						RGGELLDRILRQRYFSEREASD
						VLCTITKTMDYLHSQGVGAMP
						GAAFNILYRDESGSPESIRVCDF
				1		GFAKQLRAGNGLLMTPCYTAN
						FVAPEVSIPRLQTQACDIWSLGI
						LLYTMLAG*AVTSNRCHDTPEE
						ILARIGSGKYALSGGNWDSISD
						AAKVSTCPSVALDPHQRLTAM
		ł				QVLKHPWVVNREYLSPNQLSR
	}	1				QDVHLVKVPAG\AVRTEWWE
L	<u> </u>					WDLNRTPQAPRLEPVLSSNLAQ

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19625	49993	A	19740	144	1143	REAGRGCSAGVGGRPRRRRK GAASRARLPFSLSIMDPSLLRER ELFKKRALSTPVVEKRSASSE\S SSSS\SKKKKTKVE\HGGSSGSK QNSDHSNGSFNLKA\LSGSSG\Y KFGVLAK\IVNYMKDTGIQRGD THPLTLDEI\LDETQHLDIGLQQ KQWLMTE\ALVNNPKIEVIDGK YAFKPKYNVRDKKALLML\LD QHDQRGLGGILLEDIEEALPNS QK\AVKAL\GDQIPIC*SGPVRR KILF\F\NDKSCQFSVG*RIFRKL WEECQLVDSHGTEEKI\EEYLE ADRVFSSHARESG\PK\KVGPYS RRKGKSPAFTRKKPTFLRTSLT NHLAWNC
19626	49994	A	19741	677	1590	AHPHSPRRTLERQARISGAGRR SSPNAANRGSAPPSRRLASFCT AGSA\GRPA*SSPCSGPWTAVSA PRSPGPRGGGRPPSPGRRRPCCS VFSGGASPSSARVSSPSGSCSAS PPAAPAAWSLRGCGCWPG/VW EVHVNTGGDSV\AAPAAWGSR AACGPNLGCGSHPRAGRRAGH RCGHPWSPRPAPRPSCCGRSPR PGPPGASGCPRPAPRPPTPASRS GRPCSWRRARAAGWPRGWWG SSAGGCRRGSGRREWCPCGTA RGSPAAGVSRWTAPRCRHRRS DASASGTPARFPGAGRKGRTL WQ

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO:	sequence	nou	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	'			sequence		
10.60	1,000.5	<u> </u>	1.05.15	la a	0.170	OVA (OD LEA WAREL CHOCK) THE
19627	49995	Α	19742	754	2179	SKMSRLEAKKPSLCKSEPLTTE
						RVRTTLSVLKRIVTSCYGPSGR
						LKQLHNGFGGYVCTTSQSSALL
						SHLLVTHPILKILTASIQNHVSSF
						SDCGLFTAILCCNLIENVQRLGL
						TPTTVIRLNKHLLSLCISYLKSE
						TCGCRIPVDFSSTQILLCLVRSIL TSKPACMLTRKETEHVSALILR
						AFLLTIPENAEGHIILGKSLIVPL
						1
						KGQRVIDSTVLPGILIEMSEVQL MRLLPIKKSTALKVALFCTTLS
					:	!
-						GDTS*\PGEGTVVVSYGVSLEN
		l				AVLDQLLNLGRQLISDHVDLVL
						CQKVIHPSLKQFLNMHP\IIAIDR
						IGSDSD/VEPLDLKWTGTQPIGS
				_		LGSICPNSYGSVKDVCTAKFGS
						KHFFHLIPNEATICSLLLCNRND
						TAWDELKLTCQTALHVLQLTL
						KEPWALLGGGCTETHLAAYIR
						HKTHNDPESILKDDECTQTELQ
10600	40006	ļ.	10742		652	LIAEAFCSALESVVGSLEHDGG LMPKPGRDTTKKENFRPISLMN
19628	49996	Α	19743	1	032	
				}		INAKILNKILANQIQQHIKKLIN HDQVGFIPGMQGWFNIHKSINV
						IHHIKRTNDKNHMIISIDAGKAF
						NKIQQCFMLKTLNKLNIHGVYL
ĺ	i					KIIRAIYD/KPPANIILNGEKLEA
İ						FPLKTGIRQG/CPLLRNIV/LEVL
						ARAIRQEKEIKGIQSGNEEVKLS
						LFADDMIVYLENPIVSAQNLLK
						LISNFSKVSGYKISVQ
10620	49997	1	19744	1	834	MGDFNTPLSTLDRSTRQKVNK
19629	49997	A	19/44	1	034	DIQELNSALYQVDLIDIYRTLHP
ŀ						KSTEYTFFSAPHHTYSKIDHIVG
		1				SKALLSKCKRTEIITNCLSDHSA
						IKLELTIKKLTQNRSTTWKLNN
		1				LLLNDYWKYKQPSENKHLYAN
						KLENLEEMDKFLDTYTLPRLNQ
		}				EEVESLNRPITRSEIEAITNSLPT/
		1				KKSPGPDGFTAEFYQ\MLEVLA
		1				RAIRQEKEIKGIQLVKEEVKLSL
						FADDMIVYLENPIVSAQNLLKLI
						GNFSKVSGYKI/NVQKSQAFLY
1						TNNRQTESQIM
19630	49998	В	19745	i	1521	LIMING I ESCHAI
19631	49998	В	19746	1	1461	
17031	72777	10	12740	<u> </u>	11701	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19632	50000	A	19747	1	2347	MELKTKARELHDECTSLSSRFD
		ĺ				QLEERVSVMEDEMNEMNLPTK
		1				KSPGPDGFTAEFYQRYKEELVP
						FLLKLFQSIEKEGILPNSFYEPSII
		l				LIAKPGRDTTKKENFRPISLMNI
						NAKILNKMLANQIQQHIKKLIH
						HDQVGFIPGMQGWFNIRKSINV
	ļ					IQHINRTKDKNHMIISIDAEKAF
	1	l				DKIQQHFMLKTLNKLVLEVLA
		l				RAIRQEKEIKGIQLGKEEVKVSL
						FADDMIVYLENPTVSAQNLLKL
						IGNFSKVSGYKINVQKSQAFLY
		1				TNNRQTERQIMSELPFTIASKRI
	İ]		1		KYLGIQLTRDVKDLFKENNKPL
				ļ		LKEVKEDTNEWKNIPCSWVGRI
						NIVKMAILPKVIYRFNAIPIKLP
						MTFFTELEKTTLKFIWNQKRAC
						IAKSIFSQKNKAGGITLPDFKLY
	:					YKATVTKTAWYWYQNRDIAQ
	:					WNRTEPSEIMLHIYNYLIFDKPE
						KNKQWGKDSLFNKWCWENWL
						AICRKVKLDPFLTPYTKMNSR
						WIKDLNVRPKTIKTLEENLGITI
						QDIGVGKDFMSKTPKAMATKA
						KIDKWDLIKLKSFCTAKETTIRV
						NRQPTTWEKIFATYSSDKGLISR
						IYNELKQIYKKKTNNPIKKWAK
						DVNRHFSKEDIYAAKKHMKKC
						SSSLAIREMQIKTTMRYHLTPV
						RMAIIKKSGNNRKIQ/GGIWCD
						RIL*R*TTCRVAKEIQSL*RRI/W
	<u> </u>					KRLQRTLSIPVLDAV*PPMF*AS
19633	50001	Α	19748	1	1818	

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19634	50002	A	19749	I	1725	MELKTKARELREECRSLRSRCD QLEERPNLRLIGVPESDGENGT KLENTLQDIIQENFPNLAKQVN VQIQEIQRTPQRYSLRRATPRHII VRFTKVEMKEKMLRAAREKGR VTLKGKPIRLTADLSAETLQAR REWGPIFNILKEKNFQPRISYPA KLSFISEGEIKSFTDKQMLRDFV TTRPALQELLKEALNMERNNRS WFFEKINKIDRPLARVIKKKRE KNQIDAIKNDKEDITTNPTEIQTI IREYYKHLYPNKLENLEEMDTF LDTYTFPRLNQEEVESLNTSITG SEIVAIIS/NSIPTKKSPGPDGSTA EFYQMLEVLARAIRQEKEIKGI QLGKEEVKLSLFADDMIVYLEN PIVSAQNLLKLISNFSKVSGYKI NVQKSQAFLYTNNRQTESQIMS ELPFTIASKRIKCLGIQLTRDVK
						DLFKENYKPLLKEIKEDTNKW KNIPCSWVGRINIVKMAILPKVI YRFNAIPIKLPMTFFTDLEKTTL NFIWNQKRARITKSILSQKNKA GGITLPDFKLYYTATVTKTAW YWYQN\RWYWYQNRDIDQWN RTEPSEITPHVYNYLIF

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19635	50003	Α	19750	1	1719	MHNTDGNRFLSHWGYRQALSI
		1				SKPASASLHPSSKTKPLGTQSKT
						VVAKRNREHGKKERSSSPAME
		1				QSWMENDFDELREEGFRRSNY
		ł				SELREDIQTKGKEVENFEKNLE
]				ECITRITNTEKCLKELMELKTK
		}		<u> </u>		ARELREECRSLRSRCDQLEERR
				:		KQERSKIDTLTSQLKELEKQEQ
			-			THSKAGRRQEITKIRAELKEIET
						QKTLQKINESRSWFFERINKIDR
						PLARLIKKKREKNQIDTIKNDK
			}			GDITTDPTEIQTTIREYYKHLYA
						NKLENLEEMDTFLDTYTLPRLN
		İ				QEEVESLNRPITGSEIVAIINSLP
	1					TKKSPGPDGFTAEFYQRYKEEL
						HINRAKDKNHMIISIDAEKAFD
						KIQQPFMLKTLNKLVLEVLARA
						IRQEKEIKGIQLGKEEVKLSLFA
						DDMIVYLENPIVSAQNLLKLLS
Ì	1					NFSKVSGYKINVQKSQAFLYTN
İ	İ					NRQTESQIMSELPFTIASKRIKY
						LGIQLTRDVKHLFKENYKPLLK
						EIKEDTNKWKNIPCSWV\GRINI
						VKMAILPKVIYRFNAVPIKLPM
					:	TFFTELEKTTLKFIWNQKRACIA
						KSILSQKNKAGGITLP

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19636	50004	A	19751		2133	WRRIYQANGK*KNK/QKKAGV VILVSDKTDFKPTKIKRDKEGH YIMVKGSIQQEELTVLNIYAPN TGAPRFMKQVLRDLQRDLDPH TTIMGDFNTPLSTLDRSARQKV NKDIQELNSALHQADLINIYRIL HPKSTEYTFISAPHRTYSKIDHI VGRKALLRKYKRTEIITDCLSD HSAIKLELRIKKLTQNSSTTWK LNNLLNDYWIHNKTKAEIKM CFETSENKDTTYQNLWDTCKA VCREKFIALNAHKRKQERSKID TLTSQLKE/LEKQEQTHSKASRR KSRRNG*IPGHIHPPKTKPGRI* VPE*TNNRV*N*GNN**LTNQK KFRTRIHSQILPEHSAGSSGQG NQAGERNKGYSIRKRGSQIVPV CR*HDCIFRKPHHLSPKSP*AVK QLQQSLRIQNQRAKITSSPIHQ* QTNREPNHE*TFIHNCFKENKIP RNPTYKGCEGPIQGELQTTAQQ NKRGHKQMEEHSMLMDRKNQ YHENGHSAQGNL*IQCHPHQAT NDFLHRIGKNYFKVHMEPKKSP HCQVNPKPKEQSWRHHAT*LQ TILQGYSNQNSMVLVPKQTYRP MEKNRGLRNNTTHLRPSSL*QT *QKQEMGKGFPI**MVLGKLAS
						HM*KAETGSLPYTLYKN*FKM D*RLKC*T*NHKNLRRKPRQYH SGHRHEQGLYV*NTKSNGNKS QN*QMGSN*TKELLHSKRNYH
19637	50005	Α	19752	1	1314	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19638	50006	A	19753	1	1998	MVKGSIQQEELTILHIYAPNTG
17030	30000	l'`	17733	1	1990	APRFIKQVLSDLQRDLDSHTLI
						MGDFNTPLSTLDRSMRQKVNK
						DTQELNSALHQAALIDIYRTFH
						PKSTEYTFFSAPHHTYSKIDHIV
						GSKALLNKCKRTEIITNYLSDHS
						AIKLELRIKNLTQSRSTTWKLN
						NLLLNDYWVHNEMKAEIKMFF
		į				
		į				ETNENKDTTYQNLWDAFKA\PL
		į				ARLIKKKREKNQID/TIKNDKGD
						ITTDPTEIQTTIREYYKHLYANK
[LENLEEMDTFLDTYTLPRLNQE
ĺ						EVESLNRPITGSEIVAIINSLPTK
						KSPGPDGFTAEFYQRYKEELVP
						FLLKLFQSIEKEGILPNSFYEASII
						LIPKPGRDTTKKENFRPISLMNI
					:	DAKILNKILANRIQQHIKKLIHH
	İ	:				DQVGFIPGMQGWFNIRKSINVI
						QHINRAKDKNHMIISIDAEKAF
						DKIQQPFMLKTLNKLVLEVLAR
						AIRQEKDIKGIQLGKEEVKLSLF
						ADDMIVYLENPIVSAQNLLKLIS
						KFSKVSGYKIHVLKSQAFLYTN
						NRQTESQIMRELPFTIASKRIKY
						LGIQLTRDVKDLFKENYKPLLN
						EIKEDTNKWKNIPCSWIGRINIM
						RRSSSAPSGCHLCPPSLYLHSPN
						VELKQASFSVAGFQWKLGITAP
	1					TQPRGVGDDFVTSTSPPSLDMF
						PRLKGAENSPHRSGRD
19639	50007	Α	19754	1	3288	
19640	50008	A	19755	1264	1293	LUZZI DZI ANI VEDDE OTEGODIC
19641	50009	A	19756	264	490	VYLLIVLAVLYTNNRQTESQIM
		İ				SELPFTIASKRIKYLGIQL\TRDV
		l				KDLFKDNYIPL\LKEI*EDTSKW
10640	50010	ļ. —	10767	1	1045	KSIPCSWI
19642	50010	A	19757	1	1845	MCDENTEDICTI DECEDOMANIC
19643	50011	Α	19758	1	3144	MGDFNTPLSTLDRSSRQKVNK
						DTQELNSTLHHADLIDIYRTLHP
					}	KSTEYTFFSAPHHTYSKIDHVV
		l				GSKALLSKCKRTEIITNCLSDHS
		l				AIKPELRIKKLTQNRSTTWKLN
		1				NLLLNDYWVHNKMKAEIKMFF
						ETNENKDTTYQNLWDTFKAVS
						RGKFIALNAHKRKQKRCKIDTL
						ASQLKEVEKQEQTHSKASRRQ
						EITKIRAELKEIETQKTLQKINES
	1	I	Į.	I		RSWFLERINKIDRPLARLIKKKR
	ł	1				
19644	50012	В	19759		1743	EKNQIDVIKNDK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19645	50013	A	19760	<u> </u>	1500	MGKKQSRKTENSKNQSASPPPE
						ECSSSPAMEQSWMENDFHELR
						EEGFRSNFSELKEDVRTHSKEA
						KNLEKRLDEWLTGITSVEKSLN
						DLMELKTMARELCDKCTSFSS
			•			KFHQLEERVSVTEDQMNEMKS
		İ				WFFEKINKIDRLLARLIKKKRE
						KNQIEAIKNDKGDITTDPTEIQT
						TIREYYKHLYTNKLENLEEMDT
						FTGTYTLPRLNQEEIESLNKPIT
						GSEIEAIINSLPTKKSPGPDGFTA
						EFYQRYKEELL\FNIRKLINLIQL
						INRTQDP*PPL/VLSRIGRRVHQ
						YLENPIVSAQNLLKLKSNFSKV
						SGYKINVQKSQAFLYIKKRQTE
						SQIMSELPFTIASKRIKYLGIQLI
		l		i		RDVKDLFKENYKPLLNEIKEGT
						NKWKNIPCSWIGRINIVKMAILP
		l				KVIYRFNAIPIKLPMTFFTELEKI
						TLKFIWNQERACIAKTILSKKN
		1				KAGGIMLPDFKLYYKA\TVIKT
		1				AWDWYQNRYLDQWNRTEPSEI
		<u> </u>				TPHIYNHLIF
19646	50014	Α	19761	1	3192	
19647	50015	В	19762	180	190	
19648	50016	Α	19763	1	3212	MVKGSIQQEELTILNIYAPNTG
						ALRFIKQVLRDLQRDLDSHTIIM
					-	GDFHTPLSTLDRSTRQKVNKDI
1						QELNSALHQEDLIDIYRTLHPKS
						TEYTFFSAPHHTYSKIDHIVGSK
		l				ALLSKCKRTEIITNCLSDHSAIK
						LELRIKNLTQNRSTTWKLNNLL
						LNDYWVHNEMKAEIKMFFETN
						ENKDTTYQNLWDTFKAVCRGK
		1				FIALNAHKRKQERSKIDTLTSQL
						KELEKQEQTHSKASRRQEITKIR
	L]			AELKEIETQ

SEQ ID NO:	SEQ ID NO: of peptide sequence	ı	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	l e	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide inscrtion)
19649	50017	A	19764	1	2607	MTNHCFKRQFQKHEMLLKEDC EPGDSSFILGEQHWENMLCEPK QHPPQQGEGRAAGQAQNPLNT
						DVSPGAGAKESEEKIYQHLNYT YFKQKCMSCSGFTFSSYAMHW
						VHQAPGKGLEWVSAIGAGGGT
						YYADSVKGRFTISRDNAKNSLY
						LQMNSLRAEDMAVYYCARDTI
ŀ						DAYGDEKPSTIVTRIRVLRKLR
		İ				DTCHMGEADRTFLSPIAQHPPV
						CHVETHLGMQVANSSEEDELV
						WSKWDVDVKFILCCAKYHREQ
						LQESALDLKSLHCPSASPALFL
						ATPSQPVLTTGPGSASSQGGAE
						MGAHFLVQLVQSGAETRESFD
						PALPEMETQRTDEQCRAVHPGP
						QKERHGNMSPPLLMKGSSSPVP
						SGPGEEPSHREPVSFICGRCILLT
		1				LLLGDCLWGEGAWTQGDVLQ
						PSDRASFLAMGVNTTVQQVGD
	}					LSGYFPNSVGKACKCREFHTLT PLAHTSSTTHETFPGAQCEVQL
						VESGGGLVQPGGSLRLSCPDSG
						FTFSNHYMSWVRQAPGKGLEW
						ISYISGDSGYTNYADSVKGRFTI
						SRDNANNSPYLQMNSLRAEDT
						AVYYCVKHTARGKLEKQEQTY
}						SKASRRQEITKIRAELKETETQK
						TLQKINESRSWFFEKIYKIDRPL
						ARLIKKKREKNQIDAIKNGKGD
						ITTDPTEIQTTLREYYKHLYAN
						KLENLEGMDKFLDAYTLPRLN
19650	50018	Α	19765	1	1251	
19651	50019	Α	19766	1099	4113	
19652	50020	A	19767	1	2628	
19653	50021	В	19768	1	2970	
19654	50022	В	19769	1	4142	
19655	50023	Α	19770	1	4091	MHIVVETALSASWQNKAKPPA
						RVLLQVVPNVWFLVAVVWEL
						YPSLDLMDRSIECSSSPATEQS
						WTENDYDKLREEGFRRSNYSE
						LQEDIQTKGKDVENFEKNLEEC
						ITRITNTQKCLKELMELKTKAR
						ELREECRSLRSRCDQLEERVSV MEDEMNEMKQEGKFREKRIKR
						NEQSLQEIWDYVKRPNLRLIGV
						PESDGENGTKLKNTLQDIIQENF
						PNLARQAKVQIQEIQRMPQRYS
						LTRATPRHIIVRFTKVE
	L	L	L	L	l	LIKATIKIIIVKITKVE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19656	50024	Α	19771	156	450	LQVEIYKTVKMHCIWHKQMVS
						SYFFYQKKNELSNFEEVFLKTA
						SGMFFSTVGCHPTRCGEFEKNN
						PDLYLKELLNLAENNKGKVV\A
						IGECGLDFDRLQ
19657	50025	Α	19772	173	1009	LHGGNLQDSKDALHLAQTNG
		ŀ				MFFSTVG\IHPTRCG*/YFEKNNP
						DLYLKGVAKIFA/EKTLKGKLW
						QLGEC\GLDFD\RLQFCPKDTQL
		l				KYFEKQFE\LSEQTKLPMF\LHC
						RNSHGWNFLDIMKRNRDRCVG
		1				GVVHSFDGTKEAAS\SLIDLDLY
						IGFNG/CASPENLEANLGRFWK
		İ		1		SI\PSEKLMIEA\DAP\WCGVKST HA\GSKYIRTAFPS\KKKWESGH
		l				CLKDRNEPCHIIQILEIMSAVRD
						EDPLELANTLYNNTIKVFFSWN
		İ				IIGICLPLSIMYVKFHSKTS
19658	50026	A	19773	16	1558	QEHLQRASPRLLQLLPPLPPPPP
17030	30020	l' `	17773	1.0		PPAFGGSLSSLGSRMSRQV\VRS
		ŀ				SKFRHVFGQPAKADQ\CYEDVR
		}				VSQTT\WDSGFWC/SVNP*VLW
						ALICEASGGGAFLVLPLGKIWT
						LWDKNAPTGCGSHS/APVLDIA
						WCPHNDNVIA\SGSED\CTVMV
		İ				WEIPDGGLMLPLREPVV\TLEG
		ł				HTKRVG\IVAWHTTAQNVLLSA
			<u> </u>			GCDNVIMVWDVGP\GAAMLDT
						GPRGA/HPDANYRGDWEP\DGG
						LICTSCRDKRVRIIEPRKGTVVA
1						EKDRPHEGTRPVRAVFVSEGKI
						LTTGFSRMSERQVALWDTKHL
						EESPV/SLRKLHTRNGVPVAFFG
						P\DTNIVYLRGKG\SGPIRVFE\IT
						SE\APFLHYL\SMFS\SKESQRG\
						MGYM\PKRGRKVNK\CEIAGFY
						KLHDGR/CVSPMPKPVPRKSDL
						FQEDLGPTPPRGPDP\ALTA\EE
						WLGGRDVG\PLFISLKDGYV\PP
						KSREL\RVNRGLDP\GRRR\AAP
						EASGTPSSDAVSRLEEEMRKLQ
10650	60027	 	10774	220	475	ATVQELQKRLDRLEETVQAKL
19659	50027	Α	19774	229	475	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19660	50028	Α	19775	225	965	EWAWRSQLRSEHSRADPTRSSC
						RSCGAKMSGEENPASKPTPVQ
			,			DVQGDGRWMSLHHRFVADSK\
						DKSGSKKKNPPPPEVVFIGDSL
	İ					VQLMHQCEIWRELFSPLHAL\N
						FGIGGDGTQHVLWRLENGELE
						HIRP\KIVVVWVG/TPNNHGHTA
						EQGLLPRGQHPNPLREKNRQV
						NELGTGGTWLGHPRAHFLDAD
	1					PGFVHSDGTISHHDMYDYLHLS
						RLGYTPVCRALHSLLLRLLAQD
		1				QGQGAPLLEPAP
19661	50029	В	19776	691	1077	
19662	50030	Α	19777	481	1308	PSVAFL*PSLHRYGHTGTAGPG
						WPCAAGS*AA/SSSRRGLPSVLQ
						ASSNSSHRRCNSTPSGSAGQRR
						VRRAP*MASTTPGGTESGPCKA
						RNAFRMRSTSAFCRGQ*QRPVP
						SSSSRSSSRACA\AGETASGDP
ŀ		1				RAQRQLRS*DEESSSPSLDFRQA
						/PWPTAPCVKVSYSSSRMFSGM
						SPWRSSDNKRNSSFLRKTTGSN
						FSADLDTPPMPEVCLLPSLGYP
						LVPPVAEVHMPVPAPILPSAGIR
						RKIQKAASAKQLPLFQMDDSFS
						QHRVQGDTKK
19663	50031	С	19778	41	318	
19664	50032	Α	19779	970	1193	INNHIRDSCWTCHCSC/QPPPPPP
1						LLPQPPP*LSAPRVIERKE*DRL/
İ						CPGEQTSSPTPSPRTQSAAAAT
L						QLSLPPP
19665	50033	Α	19780 .	1	494	MDGRVMAAGGRQAPGQKKAG
						TSEAPPSSQEWPEAWYGCLQQ
						KLRAVYLVQSQPCTGPVLAPG
						AACPAAAASTPGCAQW/CGPR
						ACSPTQSSQLCAWFAFSSRCPD
						FCPNPPRPL*DPPSWPTDPADRC
						SPELTIRDSCWTCHCSCRRRHR
		L				RPCCRSRRPDSPRHG
19666	50034	C	19781	120	392	
19667	50035	Α	19782	174	375	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19668	50036	A	19783	33	1154	QFALCAGWWLCGRGGGRTPAT MSGFSTEERAVPFSLEYRVFLK NEKGQYISPFHDIPIYADKDVFH M\VVEVPRWSNAKMEIATKDP LNPIKQDVKKGKLRYVA\NLFP\ YKGYIW\NYG\AFPQTW\EDPG\ HNDKTLLGCCGLTMDP\IDVC* NWEGKVCARGEIIG\VKVLG\IL AMI\DGGGTDWKVIAINV\DDP GCQPINNDI\NDVKRLK\PGYL* KLPVDWFR\RYKVPDGKPENEF AFNAEFKDK\DFA\IDIIKSTPWT LGKALVT*ENRMGKGISCMNT NFVLRAPFKVCILDAAQSPLWD ALPTHPCGICPCHSTQQTVDKG VPITQKNLMRFPLEYQADIAYIR VPSWMVLEVKVVAFSKAFKFV
19669	50037	A	19784	1	612	EPHPNLK VTQEDDIIEEHKPRTPRPSPPLL APSPACQAFGEGHSPGHPAIST GAGPRPSASFSHPHLSGLHREPP LNWSVCPGLVWSPKPALDETH PFVQQFPHQATLEWGVFHRDV GRRWGRRLGNLQIGSEVARRR RVGGGGLQCRRLGVGASPGAD W**SG*EEACGGRWAA/GADG WGLGRPRELIGCRGWRRSLAA APGPQRPGFVSVSTPRCRPRSG METQTPALRVQQDDLFQS
19670	50038	A	19785		951	MEEGISSQIGRRENECQQKKCQ TLIKPSDLVRTHSYHENSMGETI LVIQLPPPGTAFDMWELQFKLI TQTFSHHNQLAQKTRREKRAR QEA\ERREKAERAARLAKEAKS ETSGPQIKELT\DEEAERLQLEID QKKDAENHEA\QLKNG\SLDSP GKQDTEEDEEEDEKDKGKLKP NLGNGEDLPNYRWTQTLSELD LAVPFCVNFRLKGKDMVVDIQ RRHLRVGLKGQPAII\DGELYNE VKVEESSWLIEDGKVVTVHLE KRGPADAHSRALSVLQASRLL GVHCVPGTVPNAEIQVVDFVIY FWQSVIPQASPISE

Possible nucleotide ble nucleotide insertion) RSAARLGGEQEEER IAQ\QHEGGVQELV KTDFFIGGEEGTAE IHNQL\AQKT\RREK REKAERAARLAK PQIKELTDEKA\ERL DAENHEAQLKNGS FEEDEEEEDEKDKG RADLAQ*RLNPDPC
RSAARLGGEQEEER IAQ\QHEGGVQELV KTDFFIGGEEGTAE IHNQL\AQKT\RREK RREKAERAARLAK PQIKELTDEKA\ERL DAENHEAQLKNGS IEEDEEEEDEKDKG RADLAQ*RLNPDPC
IAQ\QHEGGVQELV KTDFFIGGEEGTAE IHNQL\AQKT\RREK RREKAERAARLAK QIKELTDEKA\ERL DAENHEAQLKNGS IEEDEEEEDEKDKG RADLAQ*RLNPDPC
IAQ\QHEGGVQELV KTDFFIGGEEGTAE IHNQL\AQKT\RREK RREKAERAARLAK QIKELTDEKA\ERL DAENHEAQLKNGS IEEDEEEEDEKDKG RADLAQ*RLNPDPC
KTDFFIGGEEGTAE IHNQL\AQKT\RREK RREKAERAARLAK PQIKELTDEKA\ERL DAENHEAQLKNGS IEEDEEEEDEKDKG RADLAQ*RLNPDPC
IHNQL\AQKT\RREK RREKAERAARLAK PQIKELTDEKA\ERL DAENHEAQLKNGS FEEDEEEEDEKDKG RADLAQ*RLNPDPC
RREKAERAARLAK PQIKELTDEKA\ERL DAENHEAQLKNGS FEEDEEEEDEKDKG RADLAQ*RLNPDPC
QIKELTDEKA\ERL DAENHEAQLKNGS FEEDEEEEDEKDKG RADLAQ*RLNPDPC
DAENHEAQLKNGS FEEDEEEEDEKDKG RADLAQ*RLNPDPC
TEEDEEEEDEKDKG RADLAQ*RLNPDPC
RADLAQ*RLNPDPC
'
"ALBERT "ATTALL
F\CVNFRLKGKDMV
\VGLKGQPSDSLM
VEES\SWLIEDG\KV
NKMEWWSRLVS\S
INPENSKA\SDL\DS
KDDGMTQRTESPM
RKQEILK\KFM\DQ
KAKFPTNPCFFPP
VAMAPTIQTQAQ
SHRTLPERSGVVCR
DIPFDPKFITYPFDQ
TSLEKQHKHDLLT
DLINPDTYRIDPNVL
LEEEIQAPTSSKRSQ
V\MRKTEYISTEFN
PEVKKWGSVKQQF
DSQITAIEKTFEDA
WGEARISQHYSKP
PVFPDFKMWINPC
APKDTSGAAALEM
GMMDEEGN\QFVA
KKRKRDQEEEMD
YKIAREYNWNVK
ENYFFIFREGDGVY
RLSKRRAKA\GVQ
VKHRDMNEKELEA
ENHEPEEEEEEEME
SDEEQEKGSSSEKE
SESEREEGDRHEAS
DSSDY*ARAARDK
SEDDADSDDEDRG
DSDSGRN\GGGQR
PFPSGSEHSAQENG
SEADSDSD

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19677	50045	A	19792		1344	PGRTTRSMAEDVFLSAPIPRGC ADGRDADPTEEHMAQTERNDE EQFECQELL*CHVQVGAPEEEE EEEEDAVLVAEAEAVAAGWM LNFLCLSLCRAFREGRSEDFRRT RNSAEAIIHGLCSLTACQLRTIYI CQFLTRIAAGKTLDAQFENDER ITPLESALMIWGSIEKEHDKLHE EIQNLIKIQAIAVCMENGNFKG AEEVFERIFGDPNS\HMPFKSKL LMIISQKDTFHSFF\QHFSYNHM MEKIKSYV\NYVLSEKSSTFLM KAAAKVVESKRTRTITSQDKPS GNDVEMETEANLGYKKKC*LT NSLR*LNPQRVQYPY*GSHKNL FYLKLQHGTQQQDLNKKERRV GTPQSTKKKKESRRATESRIPVS KSQPVTPEKHRARKRQAWLWE EDKNLRSGVRKYGEGNWSKIL LHYKFNNRTSVMLKDRWRTM KKLKLISSDSED
19678	50046	A	19793		1444	IPGSTISCLKGQYPSEPFNMAED VSSAAPSPRRCADGRDP\DPTEE QMAETERNDEEQFEC\QELLEC QVQVGAPEEEEEEEEDAGLV AEAEAVAAGWMLDFLCLSLCR AFRDGRSEDFRRTRNSAKAIIY GLSMLTACPLRTIYICQ\FLTRIA AGKTP*MHRFENDE\RITPLGISP *MIWGSIEKEHDKLHEEIQNLIK IQAIAVCMENGNFKEAEEVFER IFGDPNSHMPFKSKLLMIISQKD TFHSFFQHFS\YNHMMG\KIKSY VNYVLSEKSSTFLM\KAAAKVV ESKRTRTITSQDKPSGNDVEME TEANLDTRKRSHKNLFLSKLQH GTQQQDLNKKERRVGTPQSTK/ RRKKEGRRATESRIPVSKSQP\V TPENDRARK\RQAWLWGRKDK ILRS\GVREFG\EGN\WS*ILLHY KF\NNR\TSVM\LKARW\RTMEG NLKLDFLQDSEDWIVFVKSFDG KGQFKYFDHCILFETCVIDVI
19679	50047	Α	19794	1	1020	
19680	50048	A	19795	411	769	SHLGFPGTWTRLWPSRMSPCPP QQSRNRVIQLSTSEL/APALGN WARLPQVTEQ*FRGLRKGSRT MPGSSPPATWTKGCLLA*Q*CL CPQLGWSRVAP*CRLGALFLAC SLLLAAPLRCF

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19681	50049	Α	19796	171	325	ETVAVSSAGARASSIMRIMVSS
1,,,,,,			1,7,70			RPSETSLESVFPDSAP/PGL*PPP
		l				CLSFGL
19682	50050	Α	19797	335	1536	GCSPCPPQ\QS\RNRVIQLSTSEL
						GEMELTW\Q\EIMSITEACRVLN
					:	APSEPSFEAPKPQLPYL\GPPPPT
						TY\CPC\SIHPDSGFPI\P\PPPYEL
						P\ASTSHVPDPPYSYG\NMAIPV
						SK\PLSLSGLLSEP\LQDPLALLD
	-					IGLPAGPPKPQEDPESDSGLSLN
						YSDAESLELEGTEAGRRRSELC
						RRCYPV\EYPYSLMPNSLAHSN
						YTL\PAAETPLALEPSSGP\VRA
						KPTARGE/AQGSRDERRAL\AM
						KISF\PTDKIVNFPVDDFNELLA
						RYPLTESQLALVRDIRRRGKNK
						VAAQNCRKRKLET\IVHLERKL
						ERLTNE\RERL\LRA\RGEASR\T
						LKVMRQQ\LQELY\REHFPSHLR
E .						E*NPGQTAYSP*KK*RACKQAC
		:				PNGDPSFPCGPRGT\KMEAHRL
19683	50051	Α	19798	1	408	LNLMRWEAVQARGGAFRIPRV
						DPGPTPIPHNSSSRFDCNSVSTA
						P\RVLVSGICAAGSFVLVAFSHS
						VGTSLC/GYFLLLTSPEAQDPGG
1						EEEAESAARQPLIRTEAPESKPG
						SSSSLSLRERWTVFKGLLWYIV
		<u> </u>				PLV
19684	50052	A	19799	1	759	
19685	50053	Α	19800	1	1815	
19686	50054	Α	19801	1	942	MGGCAGSRRRFSDSEGEETVPE
						PRLPLLDHQGAHWKNAVGFW
						LLGLCNNFSYVVMLSAAHDILS
						HKRTSGNQSHVDPGPTPIPHNS
						SSRFDCNSVSTAP\RVLVSGICA
						AGSFVLVAFSHSVGTSLC/GYFL
						LLTSPEAQDPGGEEEAESAARQ
						PLIRTEAPESKPGSSSSLSLRER
						WTVFKGLLWYIVPLVVVYFAE
						YFINQGLVSEGCLFRPFPFSALN
						VCEVNLVKVSQHQGGWVAVV
						TQGKKHPISCRRLNVDGMNIIS
						NDGSNKLNRSRKARVSYKVQR
			<u> </u>	<u></u>		TINASLNAATSTRAVSSNPNFV

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19687	50055	A	19802	82	1620	SLSPVADPHPSRGSPLWTLYDP
						GPSGDLNLMRWIGCAGSRRRFS
						DSEG\EETVPEPRLPLLDHEGAH
						WKNAEGFWLLGLCNNFSYVV
						MLSAAHDILSHKRTSG\NQSHV\
						DPGPNGDSPTTAYSRFDCNSVF\
						TGAVLL\ADILPTLVIKLL\APLG
						L\HLLPYRSGPRVLVSG\ICAGG
						KFRPGFAFSHSVG\TSLCGCWSF
						ASISSGLG\EVTFLSLTAFYPRA
		}				VISWWSSGTGGAGLLGALSYL
						GLTQAGLSPQQTLLSMLGIPAL
					:	LLASYFLLLTSPEAQ\DPERGEE
						AK\SQAGQPFIRTEGPE\GKPGSS
						SSLS\LRERWTVVKGLLWVPLF
						PWFEVTFPKYFITQGLFELP/LFF
						WTTSLSHAQQYRWYQMLYQA
						GVFASRSSLRS\CSIGFTW\ALAL
						LQCL\NL\VFLLADVWFGFLPSI
		ŀ				YLVF\LII\LYEGAPWGSASLREK
						PF\HNIRPWRPSDEHREVCKWR
				Ì		PPCHL*HTGGFILSGASLALPLH
	·					DFLLPSSPDTRDPQDAGHIHLW
						ARGTGYQ
19688	50056	c	19803	270	542	
19689	50057	A	19804	271	449	CLELRNLFYGACSWEVSISSDS
		1				EKPGIFNASPASAVHC**GFARP
	i			1		SSWRVLGAICTFHHSVYAKTA
	ŀ					KRKAST*RHFHH
19690	50058	A	19805	21	79	
19691	50059	Α	19806	1242	1560	WATRSSCSSMMEMPPWCPCGC
		1				AGRWRRGGASRGPREAAAGH*
		1				KPRTSNQGEQPTRRPSPYAACA
		1				RRAFAGPWASLPSPSPLSAGGL
						AWGAPCVAHPTRPSPYLLV
19692	50060	В	19807	148	2949	
19693	50061	A	19808	1335	2173	KQAHRPREEKRREEKRKEG\PS
						VPETLKKKQRNFTELKIKRL\RK
		1				KFAQQML\RKAR\RKLIYEKAK
						HYPKE\YRKMYRT*ISNGRGWA
						RK\AGQLYLVLKNPKFAV\VIRN
						QKVSMGVSP\KLRKVVVSFLRP
						SSKSSNETFVKALKQRPSD*TM
						P*GLVRGPI*LPWGYPPQS*SPV
						NEP*SYK\RGY\AKIQ*EAELPLT
						DNALIASISWVNSAFICM\DDLI
						HEIYTV\GKRFK\EANNFLWPLQ
						IVLLPRGG\MKKKTTHFVEGGD
						AGTQRSDQINRLIRRMN
19694	50062	A	19809	3	186	AO I QRODQINILLIRININ
19694	50063	A	19810	272	428	
12093	120003	14	17010	1212	720	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19696	50064	Α	19811	50	2237	RSLKAFICSILSMDKQSRDPPVK
						MQGSITTPGSIALAQAQAQAQV
j						PAKAPLAGQVSTMVTTSTTTTV
						AKTVTVTRPTGVSFKKDVPPSI
						NTTNIDTLLVATDQTERIVEPPE
						NIQEKIAFIFNNLSQSNMTQKEE
	1					LVNNTELVQSYRQQIGNVVNQ
						ANLQLFWNMYNSRRDLDINRP
						GTVPNAKTLRCPVMLVVGDNA
						PAEDGVVECNSKLDPTTTTFLK
						MADSGGLPQVTQPGKLTEAFK
						YFLQGMGYMPSASMTRLARSR
		ļ				TASLTSASSVDGSRPQACTHSE
ļ						SSEGLGQLNLKFEIEVLCKNLA
						LDINELKPGNLLKDKDRLKNLD
						EQLSAPKKDVKQPEELPPITTTT
						TSTTPATNSTCTATVPPQPQYS
Ì						YHDINVYSLAGLVPHITLNPTA
						HPQLKQCVRQAIERAVQELVHP
	İ					VVDRSIKIAMTTCEQIVRKDFA
						LDSEESRMRIAAHHMMRNLTA
						GMAMITC/RE/PLLMSISTNLKN
						SFASALRTASPLQREMMDQAA
						AQI*AQDNCFLL/CKVGGVDPK
						QLAVYEEFARNVPGFLPTNDLS
						QPTGFLAQPMKQAWATDDVA
						QIYDKCITELEQHLHAIPPTLAM
						NPQAQALRSLLEVVVLSRNSRD
						AIAALGLLQKSMENGLNYMAV
						AFAMQLVKILLVDERSVAHVT
]					EADLFHTIETLMRINAHSRGNA
						PEGMDITTRQKVNKKREGFND
19697	50065	A	19812	1	3777	KNPEFNKMVLNETYRNIKVLLT
						SDKAAANFSDRSLLKNLGHWL
	ŀ					GMITLAKNKPILHTDLDVKSLL
						LEAYVKGQQELLYVVPFVAKV
						LESSIRSVVFRPPNPWTMAIMN
		ļ				VLAELHQEHDLKLNLKFEIEVL
						CKNLALDINELKPGNLLKDKDR
						LKNLDEQLSAPKKDVKQPEELP
						PITTTTTSTTPATNTTCTATVPP
						QPQYSYHDINVYSLAGLAPHIT
						LNPTIPLFQAHPQLKQCVRQAIE
						RAVQELVHPVVDR